

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:34 ; Search time 56.5135 Seconds  
(without alignments)  
614.956 Million cell updates/sec

Title: US-09-981-876-200

Perfect score: 657

Sequence: 1 MACRCFLFLMGFLSVSQT.....PVQPEDADYVCVGVGFSP 123

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1586107 seqs, 282547505 residues

Total number of hits satisfying chosen parameters: 1586107

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : A\_Geneseq\_29Jan04:\*

1: Geneseq1980s:\*

2: Geneseq1990s:\*

3: Geneseq2000s:\*

4: Geneseq2001s:\*

5: Geneseq2002s:\*

6: Geneseq2003as:\*

7: Geneseq2003bs:\*

8: Geneseq2004s:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	657	100.0	123	2 AAW75123	AAW75123 Human sec
2	657	100.0	123	3 AAY66655	AAY66655 Membrane-
3	657	100.0	123	3 AAB24061	AAB24061 Human PRO
4	657	100.0	123	4 AAU12372	AAU12372 Human PRO
5	657	100.0	123	4 AAB65178	AAB65178 Human PRO
6	657	100.0	123	6 ABUS7993	ABUS7993 Human PRO
7	657	100.0	123	6 ABUS9071	ABUS9071 Novel hum
8	657	100.0	123	6 ABUS2583	ABUS2583 Human sec
9	657	100.0	123	6 ABO17816	ABO17816 Novel hum
10	657	100.0	123	6 ABUS60502	ABUS60502 Human sec
11	657	100.0	123	6 ABU13884	ABU13884 Human PRO
12	657	100.0	123	6 ABUS1070	ABUS1070 Human PRO
13	657	100.0	123	6 ABUS72469	ABUS72469 Novel hum
14	657	100.0	123	6 ABUS66770	ABUS66770 Human PRO
15	657	100.0	123	6 ABUS9851	ABUS9851 Novel sec
16	657	100.0	123	6 ABUS9218	ABUS9218 Human sec
17	657	100.0	123	6 ABO25915	ABO25915 Human PRO
18	657	100.0	123	6 ABO25041	ABO25041 Human sec
19	657	100.0	123	6 ABO01999	ABO01999 Novel hum
20	657	100.0	123	6 ABUS58924	ABUS58924 Human sec
21	657	100.0	123	6 ABUS2302	ABUS2302 Novel hum
22	657	100.0	123	6 ABUS9367	ABUS9367 Novel hum
23	657	100.0	123	6 ABUS67046	ABUS67046 Human sec
24	657	100.0	123	6 ABUS92133	ABUS92133 Novel hum
25	657	100.0	123	6 ABU10839	ABU10839 Human PRO

#### ALIGNMENTS

RESULT 1

AAW75123

ID AAW75123 standard; protein; 123 AA.

XX

AC AAW75123;

XX

DT 25-MAR-2003 (revised)

DT 28-JAN-1999 (first entry)

XX

DE Human secreted protein encoded by gene 67 clone HRCDF73.

XX

KW Human; secreted protein; fusion protein; gene therapy; protein therapy;  
diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia;  
developmental abnormality; foetal deficiency; blood; allergy; renal;  
immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma;  
inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;  
cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus;  
osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;  
endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.

OS Homo sapiens.

XX

XX WO9839446-A2.

PN

XX

PD 11-SEP-1998.

XX

PF 06-MAR-1998; 98WO-US004482.

XX

PR 07-MAR-1997; 97US-0038621P.

PR 07-MAR-1997; 97US-0040161P.

PR 07-MAR-1997; 97US-0040162P.

PR 07-MAR-1997; 97US-0040163P.

PR 07-MAR-1997; 97US-0040333P.

PR 07-MAR-1997; 97US-0040334P.

PR 07-MAR-1997; 97US-0040336P.

PR 07-MAR-1997; 97US-0040626P.

PR 11-APR-1997; 97US-0043311P.

PR 11-APR-1997; 97US-0043312P.

PR 11-APR-1997; 97US-0043313P.

PR 11-APR-1997; 97US-0043314P.

PR 11-APR-1997; 97US-0043315P.

PR 11-APR-1997; 97US-0043568P.

PR 11-APR-1997; 97US-0043569P.

PR 11-APR-1997; 97US-0043576P.

PR 11-APR-1997; 97US-0043578P.

PR 11-APR-1997; 97US-0043580P.

PR 11-APR-1997; 97US-0043669P.

PR 11-APR-1997; 97US-0043670P.

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PR 11-APR-1997; 97US-0043671P.
PR 11-APR-1997; 97US-0043672P.
PR 11-APR-1997; 97US-0043674P.
PR 23-MAY-1997; 97US-0047452P.
PR 23-MAY-1997; 97US-0047500P.
PR 23-MAY-1997; 97US-0047501P.
PR 23-MAY-1997; 97US-0047502P.
PR 23-MAY-1997; 97US-0047503P.
PR 23-MAY-1997; 97US-0047581P.
PR 23-MAY-1997; 97US-0047582P.
PR 23-MAY-1997; 97US-0047583P.
PR 23-MAY-1997; 97US-0047584P.
PR 23-MAY-1997; 97US-0047585P.
PR 23-MAY-1997; 97US-0047586P.
PR 23-MAY-1997; 97US-0047587P.
PR 23-MAY-1997; 97US-0047588P.
PR 23-MAY-1997; 97US-0047589P.
PR 23-MAY-1997; 97US-0047590P.
PR 23-MAY-1997; 97US-0047592P.
PR 23-MAY-1997; 97US-0047593P.
PR 23-MAY-1997; 97US-0047594P.
PR 23-MAY-1997; 97US-0047595P.
PR 23-MAY-1997; 97US-0047596P.
PR 23-MAY-1997; 97US-0047597P.
PR 23-MAY-1997; 97US-0047598P.
PR 23-MAY-1997; 97US-0047599P.
PR 23-MAY-1997; 97US-0047600P.
PR 23-MAY-1997; 97US-0047612P.
PR 23-MAY-1997; 97US-0047613P.
PR 23-MAY-1997; 97US-0047614P.
PR 23-MAY-1997; 97US-0047615P.
PR 23-MAY-1997; 97US-0047617P.
PR 23-MAY-1997; 97US-0047618P.
PR 23-MAY-1997; 97US-0047632P.
PR 23-MAY-1997; 97US-0047633P.
PR 06-JUN-1997; 97US-0048964P.
PR 06-JUN-1997; 97US-0048974P.
PR 22-AUG-1997; 97US-0056630P.
PR 22-AUG-1997; 97US-0056631P.
PR 22-AUG-1997; 97US-0056632P.
PR 22-AUG-1997; 97US-0056633P.
PR 22-AUG-1997; 97US-0056637P.
PR 22-AUG-1997; 97US-0056639P.
PR 22-AUG-1997; 97US-0056662P.
PR 22-AUG-1997; 97US-0056664P.
PR 22-AUG-1997; 97US-0056843P.
PR 22-AUG-1997; 97US-0056862P.
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PR 22-AUG-1997; 97US-0056872P.
PR 22-AUG-1997; 97US-0056874P.
PR 22-AUG-1997; 97US-0056875P.
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PR 22-AUG-1997; 97US-0056877P.
PR 22-AUG-1997; 97US-0056878P.
PR 22-AUG-1997; 97US-0056879P.
PR 22-AUG-1997; 97US-0056880P.
PR 22-AUG-1997; 97US-0056881P.
PR 22-AUG-1997; 97US-0056882P.
PR 22-AUG-1997; 97US-0056884P.
PR 22-AUG-1997; 97US-0056886P.
PR 22-AUG-1997; 97US-0056887P.
PR 22-AUG-1997; 97US-0056888P.
PR 22-AUG-1997; 97US-0056889P.
PR 22-AUG-1997; 97US-0056892P.
PR 22-AUG-1997; 97US-0056893P.
PR 22-AUG-1997; 97US-0056894P.
PR 22-AUG-1997; 97US-0056903P.
PR 22-AUG-1997; 97US-0056908P.
PR 22-AUG-1997; 97US-0056909P.
PR 22-AUG-1997; 97US-0056910P.
PR 22-AUG-1997; 97US-0056911P.
PR 05-SEP-1997; 97US-0057650P.
PR 05-SEP-1997; 97US-0057761P.

XX (HUMA-) HUMAN GENOME SCI INC.
PA Ruben SM, Rosen CA, Fischer CL, Soppet DR, Carter KC;
XX Bednarik DP, Endress GA, Yu G, Ni J, Feng P, Young PE, Greene JM;
PI Ferrie AM, Duan R, Hu J, Florence KA, Olsen HS, Ebner R, Brewer LA;
PI Moore PA, Shi Y, Lafleur DW, Li Y, Zeng Z, Kyaw H;
XX WPI: 1998-609887/51.
DR N-PSDB; AAV34220.
XX New isolated human genes and the secreted polypeptides they encode -
PT useful for diagnosis and treatment of e.g. cancers, neurological
PT disorders, immune diseases, inflammation or blood disorders.
XX Claim 1; Page 320-321; 447pp; English.
XX This sequence represents a secreted human protein encoded by the gene
CC clone detailed in the descriptor line. The gene can be used to generate
CC fusion proteins by linking to the gene to a human immunoglobulin Fc
CC portion (e.g. AAV34145) for increasing the stability of the fused protein
CC as compared to the human protein only. The invention relates to 70 novel
CC genes and their fragments (nucleic acid sequences: AAV34154-V34276; amino
CC acid sequences AA75057-W75179) which are useful for preventing, treating
CC or ameliorating medical conditions e.g. by protein or gene therapy. Also,
CC pathological conditions can be diagnosed by determining the amount of the
CC new polypeptides in a sample or by determining the presence of mutations
CC in the new polynucleotides. Specific uses are described for each of the
CC 70 polynucleotides, based on which tissues they are most highly expressed
CC in (see AAV34154 for described uses). (Updated on 25-MAR-2003 to correct
CC PF field.) (Updated on 25-MAR-2003 to correct PI field.)
XX Sequence 123 AA;
SQ
Query Match 100.0%; Score 657; DB 2; Length 123;
Best Local Similarity 100.0%; Pred. No. 4.3e-62;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MACRCLSLFLMGTFLLSVSQTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQOR 60
DB 1 MACRCLSLFLMGTFLLSVSQTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQOR 60
QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNACVLTISPQPEDDADYICSVGYG 120
DB 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNACVLTISPQPEDDADYICSVGYG 120
QY 121 FSP 123
DB 121 FSP 123
RESULT 2
AAV66655
ID AAV66655 standard; protein; 123 AA.
XX AAV66655;
XX 05-APR-2000 (first entry)
XX Membrane-bound protein PRO619.
KW Membrane-bound polypeptide; PRO polypeptide; LDL receptor; TIE ligand;
KW pharmaceutical; receptor immunoadhesin; gene mapping.
XX Homo sapiens.
XX WO9963088-A2.
XX 09-DEC-1999.
XX 02-JUN-1999; 99WO-US012252.
XX 02-JUN-1998; 98US-0087607P.
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PS Claim 12; Fig 68; 822pp; English.

XX The invention provides membrane-bound PRO polypeptides and  
 CC polynucleotides encoding them. The PRO sequences of the invention were  
 CC identified based on extracellular domain homology screening. The PRO  
 CC sequences have homology with proteins including LDL receptors, TIE  
 CC ligands and various enzymes. The membrane-bound proteins and receptor  
 CC molecules are useful as pharmaceutical and diagnostic agents. Receptor  
 CC immunoadhesins, for instance, can be used as therapeutic agents to block  
 CC receptor-ligand interactions. The membrane-bound proteins can also be  
 CC employed for screening of potential peptide or small molecule inhibitors  
 CC of the relevant receptor/ligand interaction. The PRO encoding sequences  
 CC are useful as hybridization probes, in chromosome and gene mapping and in  
 CC the generation of antisense RNA and DNA. PRO nucleic acid sequences will  
 CC also be useful for the preparation of PRO polypeptides, especially by  
 CC recombinant techniques

XX Sequence 123 AA;

Query Match 100.0%; Score 657; DB 3; Length 123;  
 Best Local Similarity 100.0%; Pred. No. 4.3e-62;  
 Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLFLLMGTFSLVSQTVLAQLDALLVFPQVQALSCITLSPQHVTIRDYGVSWYQQR 60  
 DB |||||  
 QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISPQVEDDADYYCSVGYG 120  
 DB |||||  
 QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISPQVEDDADYYCSVGYG 120  
 DB |||||  
 QY 121 FSP 123  
 DB |||||  
 QY 121 FSP 123

RESULT 3

AA024061

ID AAB24061 standard; protein; 123 AA.

AC AAB24061;

XX 29-JAN-2001 (first entry)

XX Human PRO619 protein sequence SEQ ID NO:16.

XX Human; tumour; diagnosis; neoplastic disease; neoplastic cell growth;  
 KW proliferation; tumorigenesis; identification; cancer; cytostatic;  
 KW neurotropic; neuroprotective; antiinflammatory; immunosuppressive;  
 KW immunostimulant; antiangiogenic; leukaemia; lymphoid malignancy;  
 KW neuronal disorder; glial disorder; astrocytal disorder; angiogenic;  
 KW hypothalamic disorder; glandular disorder; macrophagal disorder;  
 KW epithelial disorder; stromal disorder; macrophagal disorder;  
 KW inflammatory disorder; immunologic disorder.

XX Homo sapiens.

XX WO200053755-A2.

XX 14-SEP-2000.

XX 06-JAN-2000; 2000WO-US000376.

XX 08-MAR-1999; 99WO-US005028.

XX 02-JUN-1999; 99WO-US012252.

XX 23-JUN-1999; 99US-0141037P.

XX 07-JUL-1999; 99US-0143048P.

XX 26-JUL-1999; 99US-0145698P.

XX 30-NOV-1999; 99WO-US028313.

XX 20-DEC-1999; 99WO-US030911.

XX 05-JAN-2000; 2000WO-US000219.

XX (GETH ) GENENTECH INC.

XX Ashkenazi AJ, Baker KP, Goddard A, Gurney AL, Hillan KJ, Roy MA;  
 PI Watanabe CK, Wood WT;  
 XX WPI; 2000-572270/53.  
 DR N-PSDB; AAC58371.

XX Thirty PRO polynucleotides encoding PRO polypeptides, useful in the  
 PT treatment, diagnosis and prevention of cancer.

PS Claim 61; Fig 10; 286pp; English.

XX The present invention describes an isolated antibody that binds to one of  
 CC the human PRO proteins designated PRO212, PRO290, PRO341, PRO535, PRO619,  
 CC PRO1717, PRO809, PRO830, PRO848, PRO943, PRO1005, PRO1009, PRO1025,  
 CC PRO1030, PRO1097, PRO1107, PRO1111, PRO1153, PRO1182, PRO1184, PRO1187,  
 CC PRO1281, PRO23, PRO39, PRO834, PRO1317, PRO1710, PRO2094, PRO2145 OR  
 CC PRO2198. PRO antagonists can be used to inhibit tumour cell growth. The  
 CC PRO polypeptides and nucleotides are useful in the treatment, diagnosis  
 CC and prevention of cancer. The antibodies and other anti-tumour compounds  
 CC may be used to treat various conditions, including those characterised by  
 CC overexpression and/or activation of the amplified PRO genes. Exemplary  
 CC conditions or disorders to be treated with such antibodies and other  
 CC compounds include benign or malignant tumours (e.g. renal, liver,  
 CC kidney, bladder, breast, gastric, ovarian, colorectal, prostate,  
 CC pancreatic, lung, vulva, thyroid, hepatic carcinomas, sarcomas,  
 CC glioblastomas, and various head and neck tumours), leukaemias and  
 CC lymphoid malignancies, other disorders such as neuronal, glial,  
 CC astrocytal, hypothalamic and other glandular, macrophagal, epithelial,  
 CC stromal and blastocoele disorders, and inflammatory, angiogenic and  
 CC immunologic disorders. AAC58242 to AAC58365 represent PCR primers and  
 CC hybridisation probes used in the isolation of the human PRO sequences.  
 CC AAC58367 to AAC58396 and AAB24057 to AAB24069 represent human PRO  
 CC polynucleotide and protein sequences given in the exemplification of the  
 CC present invention

XX Sequence 123 AA;

Query Match 100.0%; Score 657; DB 3; Length 123;  
 Best Local Similarity 100.0%; Pred. No. 4.3e-62;

Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLFLLMGTFSLVSQTVLAQLDALLVFPQVQALSCITLSPQHVTIRDYGVSWYQQR 60  
 DB |||||

QY 1 MACRCLFLLMGTFSLVSQTVLAQLDALLVFPQVQALSCITLSPQHVTIRDYGVSWYQQR 60  
 DB |||||

QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISPQVEDDADYYCSVGYG 120  
 DB |||||

QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISPQVEDDADYYCSVGYG 120  
 DB |||||

QY 121 FSP 123

QY 121 FSP 123

RESULT 4

AAU12372

ID AAU12372 standard; protein; 123 AA.

XX AAU12372;

XX 24-OCT-2001 (first entry)

XX Human PRO619 polypeptide sequence.

XX Human secretory and transmembrane; PRO; mammalian; cancer; lung; breast;  
 KW prostate; cervical; tumour necrosis factor-alpha; TNF-alpha; cartilage;  
 KW ear; proliferation; glucose; free fatty acid; skeletal muscle; adipocyte;  
 KW A-peptide; factor VIIA; gene therapy.

XX Homo sapiens.

XX WO200140456-A2.





PR 20-MAR-2000; 200OWO-US007377.  
 XX (GETH ) GENENTECH INC.  
 XX Askhenazi AJ, Baker KP, Botstein D, Desnovers L, Eaton DL,  
 PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ,  
 PI Grimaldi CJ, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF,  
 PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI,  
 PI Zhang Z;  
 XX WPI; 2001-032160/04.  
 DR N-PSDB; AAF44129.  
 XX PRO polynucleotides used to produce polypeptides used to target bioactive  
 PT molecules such as toxins, radiolabels or antibodies, to specific cells,  
 PT to cause targeted cell death.  
 XX Claim 12; Fig 68; 935pp; English.  
 PS The present invention describes human secreted and transmembrane PRO  
 CC proteins. The PRO proteins have cytostatic activity. The PRO proteins can  
 CC be used for targeted delivery of bioactive molecules, such as toxins,  
 CC radiolabels or antibodies, that cause cell death. PRO nucleotide  
 CC sequences, and their fragments, can be used as hybridisation probes, in  
 CC chromosomal and gene mapping, and in the generation of anti-sense RNA and  
 CC DNA. They may also be used to produce transgenic animals which are used  
 CC to develop and screen therapeutically useful reagents. The PRO nucleotide  
 CC and protein sequence can be used for tissue typing and in treating  
 CC cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to  
 CC AAF44470 represent PCR primers and hybridisation probes used in the  
 CC isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65154 to  
 CC AAB65300 represent human PRO polynucleotide and protein sequences given  
 CC in the exemplification of the present invention  
 XX  
 SQ Sequence 123 AA;  
 Query Match 100.0%; Score 657; DB 4; Length 123;  
 Best Local Similarity 100.0%; Pred. No. 4.3e-62;  
 Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 Oy 1 MACCLSFLLMGTPFLSVQTVLAQLDALLVFPQGVQLSCTLSPOHVTIRDYGVSWYQOR 60  
 Db 1 MACCLSFLLMGTPFLSVQTVLAQLDALLVFPQGVQLSCTLSPOHVTIRDYGVSWYQOR 60  
 Oy 61 AGSAPRYLLYRSEEDHRRADIPDRFSAKDEAHNACVLTISPQVEDADYCVSGYG 120  
 Db 61 AGSAPRYLLYRSEEDHRRADIPDRFSAKDEAHNACVLTISPQVEDADYCVSGYG 120  
 Oy 121 FSP 123  
 Db 121 FSP 123  
 RESULT 6  
 ABUS7993  
 ID ABUS7993 standard; protein; 123 AA.  
 XX ABUS7993;  
 XX AC  
 XX DT 14-APR-2003 (first entry)  
 XX DE Human PRO polypeptide #25.  
 XX Human; PRO; cytostatic; tumour; cancer; breast; lung; stomach; liver;  
 KW horse; cow; dog; cat; sheep; pig; goat; rabbit; ADERT;  
 KW antibody-dependent enzyme mediated prodrug therapy.  
 XX Homo sapiens.  
 OS US2003027163-A1.  
 XX 06-FEB-2003.  
 XX

PF 15-NOV-2001; 2001US-00997666.  
 XX  
 PR 16-JUN-1997; 97US-0049787P.  
 PR 17-OCT-1997; 97US-0062250P.  
 PR 05-NOV-1997; 97WO-US020069.  
 PR 12-NOV-1997; 97US-0065186P.  
 PR 13-NOV-1997; 97US-0065311P.  
 PR 24-NOV-1997; 97US-0066770P.  
 PR 25-FEB-1998; 98US-0075945P.  
 PR 20-MAR-1998; 98US-0078910P.  
 PR 28-APR-1998; 98US-0083322P.  
 PR 07-MAY-1998; 98US-0084600P.  
 PR 28-MAY-1998; 98US-0087106P.  
 PR 02-JUN-1998; 98US-0087607P.  
 PR 02-JUN-1998; 98US-0087609P.  
 PR 02-JUN-1998; 98US-0087759P.  
 PR 03-JUN-1998; 98US-0087827P.  
 PR 04-JUN-1998; 98US-0088021P.  
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 PR 05-JUN-1998; 98US-0088167P.  
 PR 05-JUN-1998; 98US-0088202P.  
 PR 05-JUN-1998; 98US-0088212P.  
 PR 05-JUN-1998; 98US-0088217P.  
 PR 09-JUN-1998; 98US-0088655P.  
 PR 10-JUN-1998; 98US-0088734P.  
 PR 10-JUN-1998; 98US-0088738P.  
 PR 10-JUN-1998; 98US-0088742P.  
 PR 10-JUN-1998; 98US-0088810P.  
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 PR 11-JUN-1998; 98US-0088858P.  
 PR 11-JUN-1998; 98US-0088861P.  
 PR 11-JUN-1998; 98US-0088876P.  
 PR 12-JUN-1998; 98US-0089105P.  
 PR 16-JUN-1998; 98US-0089440P.  
 PR 16-JUN-1998; 98US-0089512P.  
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 PR 17-JUN-1998; 98US-0089599P.  
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 PR 25-JUN-1998; 98US-0090690P.

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PR 01-JUL-1998; 98US-0091360P.
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PR 02-JUL-1998; 98US-0091628P.
PR 02-JUL-1998; 98US-0091633P.
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Db 121 FSP 123
RESULT 7
ABUS9071
ID ABUS9071 standard; protein; 123 AA.
XX
AC ABUS9071;
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XX 28-APR-2003 (first entry)
XX
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Human; PRO; hypertrophy of neonatal heart; angiogenesis; wound healing;
cardiac insufficiency disorder; cancer; tumour; immune response;
adrenal cortical capillary endothelial growth; c-fos induction;
vascular endothelial growth factor inhibition; VEGF inhibition;
endothelial cell growth inhibitor; T-lymphocytes stimulation;
retinal neurons cell survival; rod photoreceptor cell survival;
retinal disorder; retinitis pigmentosa; kidney disorder;
mamalian kidney mesangial cell proliferation; Berger disease;
dermatitis; herpeticiformis; Crohn's disease; chondrocyte proliferation;
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chondrocyte redifferentiation; sports injury; arthritis.

XX Homo sapiens.

US2002132252-A1.

19-SEP-2002.

14-NOV-2001; 2001US-00990442.

16-JUN-1997; 97US-0049787P.

17-OCT-1997; 97US-0062250P.

05-NOV-1997; 97WO-US020069.

12-NOV-1997; 97US-0065186P.

13-NOV-1997; 97US-0065311P.

24-NOV-1997; 97US-0066770P.

25-FEB-1998; 98US-0075945P.

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07-MAY-1998; 98US-0084600P.

28-MAY-1998; 98US-0087106P.

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CC nephropathies associated with dermatitis, herpeticiformis or Crohn's  
CC disease. PRO1310, PRO844, PRO1312, PRO1192 and PRO1387 induce the  
CC proliferation and/or redifferentiation of chondrocytes in culture and are  
CC thus useful for treating sports injuries, and arthritis. This is the  
CC amino acid sequence of a novel human PRO protein

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SQ Sequence 123 AA;

Query Match 100.0%; Score 657; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 4.3e-62;  
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AC ABU82583;  
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DT 26-JUN-2003 (first entry)  
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XX  
KW Human; PRO; secreted protein; transmembrane protein;  
KW cardiac insufficiency disorders; angiogenesis; wound healing;  
KW cancerous tumour; immune response; retinal disorder; sight loss;  
KW retinitis pigmentosa; age-related macular degeneration; AMD;  
KW kidney disorder; Berger disease; nephropathy; dermatitis; herpeticiformis;  
KW Crohn's disease; sports injury; arthritis.  
XX  
OS Homo sapiens.  
XX  
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PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931936.
PR 19-DEC-2001; 2001US-00028072.
XX (GETH ) GENENTECH INC.
XX Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX WPI; 2003-341980/32.
DR N-PSDB; ACD24053.
XX New secreted and transmembrane PRO nucleic acids, for treating
PT inflammation, organ failure, atherosclerosis, cardiac injury,
PT infertility, birth defects, premature aging, acquired immunodeficiency
PT syndrome (AIDS), or cancer.
XX Claim 12; Fig 402; 660pp; English.
XX The invention describes an isolated nucleic acid (I) comprising, or which
CC has 80 % sequence identity to, or the full-length coding sequence of, one
CC of 275 nucleotide sequences, and which encodes a corresponding
CC polypeptide selected from 275 amino acid sequences, where all sequences
CC are given in the specification. The polypeptide encoded by (I) is used to
CC detect PRO polypeptides, link a bioactive molecule to a cell expressing a
CC PRO polypeptide, modulate a biological activity of a cell, stimulate the
CC release of tumour necrosis factor (TNF)-alpha from human blood, modulate
CC the uptake of glucose or free fatty acid by cells, stimulate or inhibit
CC the proliferation or differentiation of cells or gene expression,
CC stimulate the release of proteoglycans, stimulate the release of cytokine
CC from peripheral blood mononuclear cells, inhibit the binding of A-peptide
CC to factor VIIA, or detect the presence of tumour in a mammal. The nucleic
CC acid and polypeptide encoded by it, are useful for treating inflammatory
CC diseases, organ failure, atherosclerosis, cardiac injury, infertility,
CC birth defects, premature aging, acquired immunodeficiency syndrome
CC (AIDS), cancer, or diabetic complications. The nucleic acid is useful as
CC hybridisation probes, in chromosome and gene mapping, and in generating
CC antisense RNA or DNA. The polypeptides are useful as pharmaceuticals,
CC diagnostics, biosensors or bioreactors. Both are useful in tissue typing.
CC This is the amino acid sequence of a novel human secreted and
CC transmembrane PRO polypeptide
XX Sequence 123 AA;
SQ
Query Match 100.0%; Score 657; DB 6; Length 123;
Best Local Similarity 100.0%; Pred. No. 4.3e-62;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MACRCLSFLLMGTFELSVQTVLAQDLALLVFPQVLAQLSCTLSPOHVTIRDYGVSWYQQR 60
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QY 121 FSP 123
DB 121 FSP 123
RESULT 10
ABU60502
ID ABU60502 standard; protein; 123 AA.
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AC ABU60502;
XX
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DT 01-MAY-2003 (first entry)  
DE Human secreted/transmembrane protein, #43.  
XX  
XX Human; PRO; secreted; transmembrane; signal peptide; pharmaceutical;  
KW diagnostic; therapeutic; gene therapy.  
XX  
XX Homo sapiens.  
XX  
XX US2002160384-A1.  
XX PD  
XX PD 31-OCT-2002.  
XX  
XX 14-NOV-2001; 2001US-009925598.  
XX  
XX 16-JUN-1997; 97US-0049787P.  
PR 17-OCT-1997; 97US-0062250P.  
PR 05-NOV-1997; 97WO-US020069.  
PR 12-NOV-1997; 97US-0065186P.  
PR 13-NOV-1997; 97US-0065311P.  
PR 24-NOV-1997; 97US-0066770P.  
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PR 20-MAR-1998; 98US-0078910P.  
PR 28-APR-1998; 98US-0083322P.  
PR 07-MAY-1998; 98US-0084600P.  
PR 28-MAY-1998; 98US-0087106P.  
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PR 05-JUN-1998; 98US-0088655P.  
PR 10-JUN-1998; 98US-0088734P.  
PR 10-JUN-1998; 98US-0088738P.  
PR 10-JUN-1998; 98US-0088742P.  
PR 10-JUN-1998; 98US-0088810P.  
PR 10-JUN-1998; 98US-0088824P.  
PR 10-JUN-1998; 98US-0088826P.  
PR 11-JUN-1998; 98US-0088858P.  
PR 11-JUN-1998; 98US-0088861P.  
PR 11-JUN-1998; 98US-0088876P.  
PR 12-JUN-1998; 98US-0089105P.  
PR 16-JUN-1998; 98US-0089440P.  
PR 16-JUN-1998; 98US-0089512P.  
PR 16-JUN-1998; 98US-0089514P.  
PR 17-JUN-1998; 98US-0089532P.  
PR 17-JUN-1998; 98US-0089538P.  
PR 17-JUN-1998; 98US-0089598P.  
PR 17-JUN-1998; 98US-0089599P.  
PR 17-JUN-1998; 98US-0089600P.  
PR 17-JUN-1998; 98US-0089602P.  
PR 18-JUN-1998; 98US-0089901P.  
PR 18-JUN-1998; 98US-0089907P.  
PR 18-JUN-1998; 98US-0089908P.  
PR 16-SEP-1998; 98WO-US019330.  
PR 17-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 01-DEC-1998; 98WO-US021108.  
PR 05-JAN-1999; 99WO-US000106.  
PR 08-MAR-1999; 99WO-US005028.  
PR 02-JUN-1999; 99WO-US012252.  
PR 15-SEP-1999; 99WO-US021090.  
PR 15-SEP-1999; 99WO-US021547.  
PR 30-NOV-1999; 99WO-US028313.  
PR 01-DEC-1999; 99WO-US028301.  
PR 01-DEC-1999; 99WO-US028634.  
PR 16-DEC-1999; 99WO-US030095.  
PR 20-DEC-1999; 99WO-US030911.  
PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 18-FEB-2000; 2000WO-US004341.  
PR 22-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 24-FEB-2000; 2000WO-US005004.  
PR 02-MAR-2000; 2000WO-US005841.  
PR 10-MAR-2000; 2000WO-US006319.  
PR 15-MAR-2000; 2000WO-US006884.  
PR 20-MAR-2000; 2000WO-US007377.  
PR 30-MAR-2000; 2000WO-US008439.  
PR 15-MAY-2000; 2000WO-US013358.  
PR 17-MAY-2000; 2000WO-US013705.  
PR 22-MAY-2000; 2000WO-US014042.  
PR 30-MAY-2000; 2000WO-US014941.  
PR 02-JUN-2000; 2000WO-US015264.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US022031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023328.  
PR 08-NOV-2000; 2000WO-US030952.  
PR 01-DEC-2000; 2000WO-US032678.  
PR 28-FEB-2001; 2001WO-US006520.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 20-JUN-2001; 2001WO-US019692.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 28-AUG-2001; 2001US-00941992.  
XX  
XX (GETH) GENENTECH INC.  
XX  
XX Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL,  
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski P;  
PI Grimaldi JC, Gurney AL, Kljavin IG, Napier MA, Pan J, Paoni NF;  
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;  
PI Zhang Z;  
XX  
XX WPI: 2003-288106/28.  
DR N-PSDB; ABX90174.  
XX  
XX New transmembrane polypeptides and nucleic acids encoding the  
PT polypeptides, useful in gene therapy, in chromosome identification, as  
PT chromosome markers, or in generating probes.  
XX  
XX Claim 12; Fig 68; 650pp; English.  
XX  
XX The invention discloses isolated PRO secreted/transmembrane polypeptides  
CC comprising a sequence without signal peptide and the nucleic acid  
CC encoding them. The polypeptides can be used to raise antibodies that  
CC specifically bind to the PRO polypeptide, for linking a bioactive  
CC molecule to a cell expressing a PRO protein and for modulating at least  
CC one biological activity of a cell. The PRO polypeptides or  
CC polynucleotides are also useful in gene therapy, in chromosome  
CC identification, as chromosome markers, or in generating probes. The PRO  
CC polypeptides are useful as molecular markers for protein electrophoresis,  
CC and the isolated nucleic acids may be used for recombinantly expressing  
CC those markers. The PRO polypeptides and nucleic acids may also be used in  
CC tissue typing. Anti-PRO antibodies are useful in diagnostic assays for  
CC PRO, and in affinity purification of PRO from recombinant cell culture or  
CC natural sources. The sequences presented in ABU60478-ASU60624 are the PRO  
CC polynucleotides of the invention. Note: The sequence data for this patent  
CC is also available in electronic format from USPTO at  
CC seqdata.uspto.gov/sequence.html  
XX  
XX Sequence 123 AA;



Query Match 100.0%; Score 657; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 4.3e-62;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MACRCLSFLILMGTFSLVSQTVLAQLDALLVFPQVAQLSCTLSPPQHVITRDYGVSWYQQR 60  
Db 1 MACRCLSFLILMGTFSLVSQTVLAQLDALLVFPQVAQLSCTLSPPQHVITRDYGVSWYQQR 60  
QY 61 AGSAPRYLLYRSEEDHRRPADIPDRPSAAKDEAHNACVLTISFPQPEDDADYICSVGYG 120  
Db 61 AGSAPRYLLYRSEEDHRRPADIPDRPSAAKDEAHNACVLTISFPQPEDDADYICSVGYG 120  
QY 121 FSP 123  
Db 121 FSP 123

RESULT 11

ABU13884  
ID ABU13884 standard; protein; 123 AA.

XX AC ABU13884;

XX XX 26-FEB-2003 (first entry)

XX DE Human PRO619 polypeptide.

XX KW Human: PRO polypeptide; secreted protein; transmembrane protein;  
XX KW genetic disorder; antibacterial; immunosuppressive.

XX OS Homo sapiens.

XX PN US2002103125-A1.

XX PD 01-AUG-2002.

XX PF 20-NOV-2001; 2001US-00989731.

XX PR 16-JUN-1997; 97US-0049787P.

XX PR 17-OCT-1997; 97US-0062250P.

XX PR 05-NOV-1997; 97WO-US020069.

XX PR 12-NOV-1997; 97US-0065186P.

XX PR 13-NOV-1997; 97US-0065311P.

XX PR 24-NOV-1997; 97US-0066770P.

XX PR 25-FEB-1998; 98US-0075945P.

XX PR 20-MAR-1998; 98US-0078910P.

XX PR 28-APR-1998; 98US-0083322P.

XX PR 07-MAY-1998; 98US-0084600P.

XX PR 28-MAY-1998; 98US-0087106P.

XX PR 02-JUN-1998; 98US-0087607P.

XX PR 02-JUN-1998; 98US-0087609P.

XX PR 02-JUN-1998; 98US-0087759P.

XX PR 03-JUN-1998; 98US-0087827P.

XX PR 04-JUN-1998; 98US-0088021P.

XX PR 04-JUN-1998; 98US-0088035P.

XX PR 04-JUN-1998; 98US-0088048P.

XX PR 04-JUN-1998; 98US-0088080P.

XX PR 04-JUN-1998; 98US-0088028P.

XX PR 04-JUN-1998; 98US-0088029P.

XX PR 04-JUN-1998; 98US-0088030P.

XX PR 04-JUN-1998; 98US-0088033P.

XX PR 05-JUN-1998; 98US-0088167P.

XX PR 05-JUN-1998; 98US-0088202P.

XX PR 05-JUN-1998; 98US-0088212P.

XX PR 05-JUN-1998; 98US-0088217P.

XX PR 09-JUN-1998; 98US-0088655P.

XX PR 10-JUN-1998; 98US-0088734P.

XX PR 10-JUN-1998; 98US-0088738P.

XX PR 10-JUN-1998; 98US-0088742P.

XX PR 10-JUN-1998; 98US-0088810P.

XX PR 10-JUN-1998; 98US-0088824P.

XX PR 10-JUN-1998; 98US-0088826P.

XX PR 11-JUN-1998; 98US-0088858P.

PR 11-JUN-1998; 98US-0088861P.  
PR 11-JUN-1998; 98US-0088876P.  
PR 12-JUN-1998; 98US-0089105P.  
PR 16-JUN-1998; 98US-0089440P.  
PR 16-JUN-1998; 98US-0089512P.  
PR 16-JUN-1998; 98US-0089514P.  
PR 17-JUN-1998; 98US-0089532P.  
PR 17-JUN-1998; 98US-0089538P.  
PR 17-JUN-1998; 98US-0089588P.  
PR 17-JUN-1998; 98US-0089599P.  
PR 17-JUN-1998; 98US-0089600P.  
PR 17-JUN-1998; 98US-0089653P.  
PR 18-JUN-1998; 98US-0089801P.  
PR 18-JUN-1998; 98US-0089907P.  
PR 18-JUN-1998; 98US-0089908P.  
PR 16-SEP-1998; 98WO-US019330.  
PR 16-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 01-DEC-1998; 98WO-US025108.  
PR 05-JAN-1999; 99WO-US000106.  
PR 08-MAR-1999; 99WO-US005028.  
PR 02-JUN-1999; 99WO-US012252.  
PR 15-SEP-1999; 99WO-US021090.  
PR 15-SEP-1999; 99WO-US021547.  
PR 30-NOV-1999; 99WO-US028313.  
PR 01-DEC-1999; 99WO-US028301.  
PR 16-DEC-1999; 99WO-US028634.  
PR 20-DEC-1999; 99WO-US030095.  
PR 06-JAN-2000; 99WO-US030911.  
PR 06-JAN-2000; 2000WO-US000219.  
PR 11-FEB-2000; 2000WO-US000376.  
PR 18-FEB-2000; 2000WO-US003565.  
PR 22-FEB-2000; 2000WO-US004341.  
PR 24-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 24-FEB-2000; 2000WO-US005004.  
PR 02-MAR-2000; 2000WO-US005841.  
PR 10-MAR-2000; 2000WO-US006319.  
PR 15-MAR-2000; 2000WO-US006884.  
PR 20-MAR-2000; 2000WO-US007377.  
PR 30-MAR-2000; 2000WO-US008439.  
PR 15-MAY-2000; 2000WO-US013358.  
PR 17-MAY-2000; 2000WO-US013705.  
PR 22-MAY-2000; 2000WO-US014042.  
PR 30-MAY-2000; 2000WO-US014941.  
PR 02-JUN-2000; 2000WO-US015264.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US02031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023328.  
PR 08-NOV-2000; 2000WO-US030952.  
PR 01-DEC-2000; 2000WO-US032678.  
PR 28-FEB-2001; 2001WO-US006520.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 20-JUN-2001; 2001WO-US019692.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 28-AUG-2001; 2001US-00941992.

(GETH ) GENENTECH LTD.

XX Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;  
XX Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;  
XX Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;  
XX Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams FM, Wood WI;  
XX Zhang Z;

XX WPI; 2003-102117/09.

XX N-PSDB; ABX64020.

XX Novel secreted and transmembrane polypeptide for modulating biological  
XX activity of cell expressing the polypeptide, identifying agonists or  
XX antagonists of polypeptide, and as molecular weight markers.

XX Claim 12; Fig 68; 649pp; English.

XX The present invention relates to the isolation of novel human PRO

PS polypeptides, and the polynucleotide sequences encoding them. The PRO

PS polypeptides are secreted and transmembrane proteins. The PRO

CC polypeptides are useful for detecting other PRO polypeptides, for linking

CC polypeptides are useful for detecting other PRO polypeptides, for modulating

CC polypeptides are useful for detecting other PRO polypeptides, and for for

CC biological activities of cells expressing PRO polypeptides, and for for

CC identifying agonists or antagonists. The polynucleotide sequences

CC encoding PRO polypeptides are useful as hybridisation probes, in

CC chromosome and gene mapping, in the generation of antisense RNA and DNA,

CC in the preparation of PRO polypeptides, for generating transgenic animals

CC or knockout animals, to construct hybridisation probes for mapping the

CC gene which encodes the PRO polypeptide, and for the genetic analysis of

CC individuals with genetic disorders, in gene therapy, for chromosome

CC identification, as chromosome markers, and for generating probes for PCR,

CC Northern analysis, Southern analysis and Western analysis. ABU13860-

CC ABU14006 represent the human PRO polypeptides of the invention. Note: The

CC sequence data for this patent was obtained in electronic format directly

CC from the USPTO web site at [seqdata.uspto.gov/psipsdidentry.html](http://seqdata.uspto.gov/psipsdidentry.html)

XX SQ Sequence 123 AA;

Query Match 100.0%; Score 657; DB 6; Length 123;

Best Local Similarity 100.0%; Pred. No. 4,3e-62;

Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRLGFLMGTLFSLVSQTVLAQDALLVFPFGVAQLSCTLSFQHVITRDYGVNQOR 60

DB 1 MACRLGFLMGTLFSLVSQTVLAQDALLVFPFGVAQLSCTLSFQHVITRDYGVNQOR 60

QY 61 AGSAPRYLLYRSEEDHHRPADIPDRFSAKDEAHNAACVLTISFVQPEDDADYVCVGYG 120

DB 61 AGSAPRYLLYRSEEDHHRPADIPDRFSAKDEAHNAACVLTISFVQPEDDADYVCVGYG 120

QY 121 FSP 123.

DB 121 FSP 123

RESULT 12

ID ABUS1070 standard; protein; 123 AA.

AC ABUS1070;

XX 23-JUN-2003 (first entry)

XX Human PRO polypeptide #201.

XX Human; PRO polypeptide; secreted and transmembrane protein;

XX anti-PRO antibody; diagnostic assay; gene expression; diabetes;

XX bone disorder; cartilage disorder; rheumatoid arthritis; obesity;

XX sports injury; osteoarthritis; hyper-insulinaemia; hypo-insulinaemia;

XX hearing loss; coagulation disorder; stroke; heart attack; cardiac;

XX antidiabetic; anorectic; vulnery; antiarthritic; osteopathic;

XX antirheumatic; auditory; cerebroprotective; angiogenic.

OS Homo sapiens.

XX US2003004311-A1.

XX 02-JAN-2003.

XX 19-DEC-2001; 2001US-00028072.

XX 18-JUN-1997; 97US-0049911P.

XX 26-AUG-1997; 97US-0056974P.

XX 17-SEP-1997; 97US-00591113P.

XX 17-SEP-1997; 97US-00591113P.

XX 17-SEP-1997; 97US-00591117P.

XX 17-SEP-1997; 97US-0059122P.

17-SEP-1997; 97US-0059184P.

18-SEP-1997; 97US-0053263P.

19-SEP-1997; 97US-0053252P.

19-SEP-1997; 97US-0053588P.

17-SEP-1997; 97US-0059836P.

17-OCT-1997; 97US-0062250P.

17-OCT-1997; 97US-0062285P.

17-OCT-1997; 97US-0062287P.

17-OCT-1997; 97US-0063755P.

24-OCT-1997; 97US-0062814P.

24-OCT-1997; 97US-0062818P.

24-OCT-1997; 97US-0063045P.

24-OCT-1997; 97US-0063082P.

24-OCT-1997; 97US-0063127P.

27-OCT-1997; 97US-0063327P.

27-OCT-1997; 97US-0063329P.

28-OCT-1997; 97US-0063550P.

28-OCT-1997; 97US-0063561P.

29-OCT-1997; 97US-0063704P.

29-OCT-1997; 97US-0063733P.

29-OCT-1997; 97US-0063735P.

29-OCT-1997; 97US-0063738P.

03-NOV-1997; 97US-0064248P.

07-NOV-1997; 97US-0064809P.

12-NOV-1997; 97US-0065186P.

17-NOV-1997; 97US-0065846P.

21-NOV-1997; 97US-0066364P.

24-NOV-1997; 97US-0066453P.

24-NOV-1997; 97US-0066511P.

24-NOV-1997; 97US-0066770P.

11-DEC-1997; 97US-0069212P.

11-DEC-1997; 97US-0069278P.

11-DEC-1997; 97US-0069334P.

16-DEC-1997; 97US-0069694P.

23-JAN-1998; 98US-0072320P.

04-FEB-1998; 98US-0073612P.

09-FEB-1998; 98US-0074086P.

09-FEB-1998; 98US-0074092P.

12-MAR-1998; 98US-0077911P.

20-MAR-1998; 98US-0078910P.

25-MAR-1998; 98US-0079294P.

27-MAR-1998; 98US-0079663P.

27-MAR-1998; 98US-0079728P.

31-MAR-1998; 98US-0080185P.

12-JUN-1998; 98WO-US012456.

14-JUL-1998; 98WO-US014552.

28-AUG-1998; 98WO-US017888.

10-SEP-1998; 98WO-US018824.

14-SEP-1998; 98WO-US019093.

14-SEP-1998; 98WO-US019094.

14-SEP-1998; 98WO-US019177.

16-SEP-1998; 98WO-US019330.

17-SEP-1998; 98WO-US019437.

27-OCT-1998; 98WO-US021141.

29-OCT-1998; 98WO-US022991.

29-OCT-1998; 98WO-US022992.

20-NOV-1998; 98WO-US024855.

01-DEC-1998; 98WO-US025108.

05-JAN-1999; 99WO-US000106.

08-MAR-1999; 99WO-US005028.

10-MAR-1999; 99WO-US005190.

20-APR-1999; 99WO-US008615.

14-MAY-1999; 99WO-US010733.

02-JUN-1999; 99WO-US012252.

01-SEP-1999; 99WO-US020111.

08-SEP-1999; 99WO-US020594.

13-SEP-1999; 99WO-US020944.

15-SEP-1999; 99WO-US021090.

15-SEP-1999; 99WO-US021547.

29-OCT-1999; 99WO-US023089.

29-NOV-1999; 99WO-US028214.

30-NOV-1999; 99WO-US028313.

30-NOV-1999; 99WO-US028409.

PR 01-DEC-1999; 99WO-US028301.  
PR 01-DEC-1999; 99WO-US028634.  
PR 02-DEC-1999; 99WO-US028551.  
PR 02-DEC-1999; 99WO-US028564.  
PR 02-DEC-1999; 99WO-US028565.  
PR 16-DEC-1999; 99WO-US030095.  
PR 20-DEC-1999; 99WO-US030311.  
PR 20-DEC-1999; 99WO-US030399.  
PR 30-DEC-1999; 99WO-US031243.  
PR 30-DEC-1999; 99WO-US031274.  
PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000277.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 18-FEB-2000; 2000WO-US004341.  
PR 18-FEB-2000; 2000WO-US004342.  
PR 22-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 24-FEB-2000; 2000WO-US005004.  
PR 01-MAR-2000; 2000WO-US005501.  
PR 02-MAR-2000; 2000WO-US005746.  
XX (GETH ) GENENTECH INC.  
PA Baker KP, Beresini M, DeForge L, Desnoyers L, Filvaroff E, Gao W;  
XX Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;  
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;  
XX  
DR WPI; 2003-352836/33.  
DR N-ESDE; ACA67194.  
XX  
XX New isolated PRO polypeptide useful for treating diabetes, rheumatoid  
PT arthritis, sports injuries, obesity, hearing loss in mammals, stroke, or  
PT heart attack.  
XX  
XX Claim 12; Fig 402; 643pp; English.  
XX  
XX The present invention relates to the isolation of novel human PRO  
CC polypeptides, and the polynucleotide sequences encoding them. The PRO  
CC polypeptides are secreted and transmembrane proteins. The PRO  
CC polypeptides and polynucleotides are useful for preparing a medicament  
CC useful in the treatment of diabetes, bone and/or cartilage disorders  
CC (e.g. rheumatoid arthritis, sports injuries, osteoarthritis), obesity,  
CC hyper- or hypo-insulinaemia, hearing loss, and coagulation disorders  
CC (e.g. stroke, heart attack). Anti-PRO antibodies are useful in diagnostic  
CC assays for PRO, by detecting its expression in specific cells, tissues or  
CC serum, and for affinity purification of PRO from recombinant cell culture  
CC or natural sources. ABUS0870-ABUS1144 represent the human PRO  
CC polypeptides of the invention. Note: The sequence data for this patent  
CC was obtained in electronic format directly from the USPTO web site at  
CC seqdata.uspto.gov/psipsdEntry.html  
XX  
SQ Sequence 123 AA;  
  
Query Match 100.0%; Score 657; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 4.3e-62;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
Qy 1 MACRCLSFLLMGTLFSLVSQTFLAQDLALLVPPGVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
Db 1 MACRCLSFLLMGTLFSLVSQTFLAQDLALLVPPGVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
  
Qy 61 AGSAPRYLLYRSDEHHRPADIPDRFSAKDEAHNACVLTI SPVQPEDDADYCVSYG 120  
Db 61 AGSAPRYLLYRSDEHHRPADIPDRFSAKDEAHNACVLTI SPVQPEDDADYCVSYG 120  
  
Qy 121 FSP 123  
Db 121 FSP 123  
  
RESULT 13  
ABU72469

ID ABU72469 standard; protein; 123 AA.  
XX  
AC ABU72469;  
XX  
XX  
DT 17-JUN-2003 (first entry)  
XX  
DE Novel human secreted and transmembrane protein PRO619.  
XX  
XX Human; secreted and transmembrane protein; cytostatic; anti-HIV;  
KW virucide; hepatotropic; antiinflammatory; neuroprotective; gene therapy;  
KW PRO; pharmaceutical; diagnostic; biosensor; bioreactor; malignancy;  
KW cancer; ovarian cancer; colorectal cancer; Kaposi's sarcoma; leukaemia;  
KW lymphoma; hepatitis B; multiple sclerosis; Crohn's disease;  
KW drug screening.  
XX  
XX Homo sapiens.  
XX US2003003531-A1.  
XX  
XX 02-JAN-2003.  
XX  
XX 19-NOV-2001; 2001US-00889734.  
XX  
XX 16-JUN-1997; 97US-0049787P.  
PR 17-OCT-1997; 97US-0062250P.  
PR 05-NOV-1997; 97WO-US020069.  
PR 12-NOV-1997; 97US-0065186P.  
PR 13-NOV-1997; 97US-0065311P.  
PR 24-NOV-1997; 97US-0068770P.  
PR 25-FEB-1998; 98US-0075945P.  
PR 20-MAR-1998; 98US-0078910P.  
PR 28-APR-1998; 98US-0083322P.  
PR 07-MAY-1998; 98US-0084600P.  
PR 28-MAY-1998; 98US-0087106P.  
PR 02-JUN-1998; 98US-0087607P.  
PR 02-JUN-1998; 98US-0087609P.  
PR 02-JUN-1998; 98US-0087759P.  
PR 03-JUN-1998; 98US-0087827P.  
PR 04-JUN-1998; 98US-0088021P.  
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PR 04-JUN-1998; 98US-0088033P.  
PR 04-JUN-1998; 98US-0088326P.  
PR 05-JUN-1998; 98US-0088167P.  
PR 05-JUN-1998; 98US-0088202P.  
PR 05-JUN-1998; 98US-0088212P.  
PR 05-JUN-1998; 98US-0088217P.  
PR 09-JUN-1998; 98US-0088655P.  
PR 10-JUN-1998; 98US-0088734P.  
PR 10-JUN-1998; 98US-0088738P.  
PR 10-JUN-1998; 98US-0088742P.  
PR 10-JUN-1998; 98US-0088810P.  
PR 10-JUN-1998; 98US-0088824P.  
PR 10-JUN-1998; 98US-0088826P.  
PR 11-JUN-1998; 98US-0088858P.  
PR 11-JUN-1998; 98US-0088861P.  
PR 12-JUN-1998; 98US-0089105P.  
PR 12-JUN-1998; 98US-0089440P.  
PR 16-JUN-1998; 98US-0089512P.  
PR 16-JUN-1998; 98US-0089514P.  
PR 17-JUN-1998; 98US-0089532P.  
PR 17-JUN-1998; 98US-0089538P.  
PR 17-JUN-1998; 98US-0089598P.  
PR 17-JUN-1998; 98US-0089599P.  
PR 17-JUN-1998; 98US-0089600P.  
PR 17-JUN-1998; 98US-0089653P.  
PR 18-JUN-1998; 98US-0089801P.  
PR 18-JUN-1998; 98US-0089907P.  
PR 18-JUN-1998; 98US-0089908P.

PR 16-SEP-1998; 98WO-US019330.  
PR 17-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 01-DEC-1998; 98WO-US025108.  
PR 05-JAN-1999; 98WO-US000106.  
PR 08-MAR-1999; 98WO-US005028.  
PR 02-JUN-1999; 98WO-US012252.  
PR 15-SEP-1999; 98WO-US021090.  
PR 15-SEP-1999; 98WO-US021547.  
PR 30-NOV-1999; 98WO-US028313.  
PR 01-DEC-1999; 98WO-US028301.  
PR 01-DEC-1999; 98WO-US028341.  
PR 16-DEC-1999; 98WO-US030095.  
PR 20-DEC-1999; 98WO-US030911.  
PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 18-FEB-2000; 2000WO-US004341.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 24-FEB-2000; 2000WO-US005004.  
PR 02-MAR-2000; 2000WO-US005841.  
PR 10-MAR-2000; 2000WO-US006319.  
PR 15-MAR-2000; 2000WO-US006884.  
PR 20-MAR-2000; 2000WO-US007377.  
PR 30-MAR-2000; 2000WO-US008439.  
PR 15-MAY-2000; 2000WO-US013358.  
PR 17-MAY-2000; 2000WO-US013705.  
PR 22-MAY-2000; 2000WO-US014042.  
PR 30-MAY-2000; 2000WO-US014941.  
PR 02-JUN-2000; 2000WO-US015264.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US022031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023328.  
PR 08-NOV-2000; 2000WO-US030952.  
PR 01-DEC-2000; 2000WO-US032678.  
PR 28-FEB-2001; 2001WO-US006520.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 20-JUN-2001; 2001WO-US019692.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 28-AUG-2001; 2001WO-US021992.  
PA (GETH ) GENENTECH INC.  
XX  
PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;  
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;  
PI Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;  
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;  
PI Zhang Z;  
XX  
DR WPI; 2003-352829/33.  
DR N-PSDB; ACA64242.  
XX  
PT New genes and secreted and transmembrane polypeptides (e.g. PRO183 or  
PT PRO184), useful for treating or diagnosing e.g. ovarian cancer, Kaposi's  
PT sarcoma, leukemia, lymphoma, hepatitis B, multiple sclerosis or Crohn's  
PT disease.  
XX  
FS Claim 12; Fig 68; 663pp; English.  
XX  
CC The invention describes a new isolated nucleic acid molecule comprising  
CC the full length coding sequence of the DNA deposited with the American  
CC Type Culture Collection (e.g. ATCC Deposit No. 209621, 552-PTA, 819-PTA,  
CC 209439, 203135, etc.) or a sequence with at least 80% identity to a DNA  
CC encoding a PRO polypeptide. The PRO polypeptides or polynucleotides are  
CC useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These  
CC are particularly useful for detecting or treating e.g. malignancies or  
CC cancers (e.g. ovarian cancer, colorectal cancer, Kaposi's sarcoma,  
CC leukaemia or lymphoma), hepatitis B, multiple sclerosis, or Crohn's  
CC disease in mammals. The PRO polypeptides are useful in drug screening,  
CC particularly as targets for therapeutic intervention in these diseases,  
XX

CC and in the diagnostic determination of the presence of these diseases.  
CC The PRO polypeptides are also useful as molecular weight markers, or for  
CC chromosome identification. The PRO genes are useful as hybridisation  
CC probes, or for screening libraries of human cDNA, genomic DNA or mRNA.  
CC The PRO genes may also be used in gene therapy, particularly for  
CC replacing a defective gene. This is the amino acid sequence of a novel  
CC human secreted and transmembrane PRO polypeptide  
XX  
SQ Sequence 123 AA;

Query Match 100.0%; Score 657; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 4.3e-6;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLSFLMGTFLSVSQTFLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
|||||  
DB 1 MACRCLSFLMGTFLSVSQTFLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
|||||  
QY 61 AGSAPRYLLYRSEEDHHRPADIPDRFSAKDEAHNACVLITISPVQEDDADYCSVGYG 120  
|||||  
DB 61 AGSAPRYLLYRSEEDHHRPADIPDRFSAKDEAHNACVLITISPVQEDDADYCSVGYG 120  
|||||  
QY 121 RSP 123  
|||||  
DB 121 RSP 123  
|||||

## RESULT 14

ABU66770  
ID ABU66770 standard; protein; 123 AA.

AC ABU66770;

XX 23-MAY-2003 (first entry)

XX Human PRO polypeptide #201.

XX Human; PRO polypeptide; secreted and transmembrane protein;  
XX tumour necrosis factor-alpha; TNF-alpha; blood; proliferation;  
XX differentiation; chondrocyte; tumour; Genetic disorder; cytostatic.

OS Homo sapiens.

XX US2003036180-A1.

XX 20-FEB-2003.

XX 09-MAY-2002; 2002US-00143114.

XX 31-MAR-1997; 97WO-US005230.

XX 12-JUN-1998; 98WO-US012456.

XX 14-JUL-1998; 98WO-US014552.

XX 28-AUG-1998; 98WO-US017888.

XX 10-SEP-1998; 98WO-US018824.

XX 14-SEP-1998; 98WO-US019093.

XX 14-SEP-1998; 98WO-US019094.

XX 16-SEP-1998; 98WO-US019177.

XX 17-SEP-1998; 98WO-US019330.

XX 07-OCT-1998; 98WO-US021947.

XX 29-OCT-1998; 98WO-US022991.

XX 29-OCT-1998; 98WO-US022992.

XX 20-NOV-1998; 98WO-US024855.

XX 01-DEC-1998; 98WO-US025108.

XX 08-JAN-1999; 99WO-US000106.

XX 08-MAR-1999; 99WO-US005028.

XX 10-MAR-1999; 99WO-US005390.

XX 20-APR-1999; 99WO-US008615.

XX 14-MAY-1999; 99WO-US010733.

XX 02-JUN-1999; 99WO-US012252.

XX 01-SEP-1999; 99WO-US020111.

XX 08-SEP-1999; 99WO-US020594.

XX 13-SEP-1999; 99WO-US020944.

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PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 29-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US0003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006684.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032578.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 01-MAR-2001; 2001WO-US006520.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00806889.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 05-JUN-2001; 2001WO-US017800.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019892.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.

PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX (GETH ) GENENTECH INC.
PI Baker KP, Beresini M, DeForge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX WPI; 2003-332040/31.
DR N-PSDB; ACA03803.
XX
XX New secreted and transmembrane PRO nucleic acids, useful for gene
PT therapy, in chromosome and gene mapping, as chromosome markers, in tissue
PT typing, and in chromosome identification.
XX
XX Claim 12; Fig 402; 660pp; English.
XX
XX The present invention relates to the isolation of novel human PRO
CC polypeptides, and the polynucleotide sequences encoding them. The PRO
CC polypeptides are secreted and transmembrane proteins. The PRO
CC polypeptides are useful for detecting other PRO polypeptides, for linking
CC bioactive molecules to cells expressing PRO polypeptides, for modulating
CC biological activities of cells expressing PRO polypeptides, and for
CC identifying agonists or antagonists. The PRO polypeptides are useful for
CC for stimulating the release of tumour necrosis factor (TNF)-alpha from
CC human blood, for stimulating the proliferation or differentiation of
CC chondrocytes, and detecting the presence of tumours. The polynucleotide
CC sequences encoding PRO polypeptides are useful as hybridisation probes,
CC in chromosome and gene mapping, in the generation of antisense RNA and
CC DNA, in the preparation of PRO polypeptides, for generating transgenic
CC animals or knockout animals, for the genetic analysis of individuals with
CC genetic disorders, and in gene therapy. ABU6570-ABU66844 represent the
CC human PRO polypeptides of the invention. Note: The sequence data for this
CC patent was obtained in electronic format directly from the USPTO web site
CC at seqdata.uspto.gov/psipspidentry.html
XX
XX Sequence 123 AA;
SQ
Query Match 100.0%; Score 657; DB 6; Length 123;
Best Local Similarity 100.0%; Pred. No. 4.3e-62;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MACRCISFLLMGTFLSVSQTVLAQLDALLVFPQGVAQLSCTLSPOHQVTRIDYGVSWYQQR 60
Db 1 MACRCISFLLMGTFLSVSQTVLAQLDALLVFPQGVAQLSCTLSPOHQVTRIDYGVSWYQQR 60
QY 61 AGSAPRYLLYRSBEDHHRPADIPDRFSAAKDEAHNACVLITSPVQPEDDADYCSVYG 120
Db 61 AGSAPRYLLYRSBEDHHRPADIPDRFSAAKDEAHNACVLITSPVQPEDDADYCSVYG 120
QY 121 FSP 123
Db 121 FSP 123
RESULT 15
ABU59851
XX ID ABU59851 standard; protein; 123 AA.
XX AC ABU59851;
XX
XX 13-MAY-2003 (first entry)
XX
XX Novel secreted and transmembrane protein PRO619.
XX Human; PRO; hypertrophy of neonatal heart; angiogenesis; wound healing;
KW cardiac insufficiency disorder; cancer; tumour; immune response;
KW adrenal cortical capillary endothelial growth; c-fos induction;
KW vascular endothelial growth factor inhibition; VEGF inhibition;
KW endothelial cell growth inhibitor; T-lymphocytes stimulation;
KW retinal neurons cell survival; rod photoreceptor cell survival;
KW retinal disorder; retinitis pigmentosa; kidney disorder;
```

KW mammalian kidney mesangial cell proliferation; Berger disease;  
KW dermatitis; herpeticiformis; Crohn's disease; chondrocyte proliferation;  
KW chondrocyte redifferentiation; sports injury; arthritis.  
XX Homo sapiens.  
OS US2003017563-A1.  
PN 23-JAN-2003.  
PD 07-MAY-2002; 2002US-00140808.  
XX 31-MAR-1997; 97WO-US005230.  
XX 12-JUN-1998; 98WO-US012456.  
PR 28-AUG-1998; 98WO-US014552.  
PR 10-SEP-1998; 98WO-US018824.  
PR 14-SEP-1998; 98WO-US019093.  
PR 14-SEP-1998; 98WO-US019094.  
PR 14-SEP-1998; 98WO-US019177.  
PR 16-SEP-1998; 98WO-US019330.  
PR 17-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 29-OCT-1998; 98WO-US022991.  
PR 29-OCT-1998; 98WO-US022992.  
PR 20-NOV-1998; 98WO-US024855.  
PR 01-DEC-1998; 98WO-US025108.  
PR 05-JAN-1999; 99WO-US000106.  
PR 08-MAR-1999; 99WO-US005028.  
PR 10-MAR-1999; 99WO-US005190.  
PR 20-APR-1999; 99WO-US008615.  
PR 14-MAY-1999; 99WO-US010733.  
PR 02-JUN-1999; 99WO-US012252.  
PR 01-SEP-1999; 99WO-US020111.  
PR 08-SEP-1999; 99WO-US020594.  
PR 13-SEP-1999; 99WO-US020944.  
PR 15-SEP-1999; 99WO-US021090.  
PR 05-OCT-1999; 99WO-US021547.  
PR 29-NOV-1999; 99WO-US028214.  
PR 30-NOV-1999; 99WO-US028313.  
PR 30-NOV-1999; 99WO-US028409.  
PR 01-DEC-1999; 99WO-US028401.  
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PR 02-DEC-1999; 99WO-US028564.  
PR 02-DEC-1999; 99WO-US028565.  
PR 16-DEC-1999; 99WO-US030395.  
PR 20-DEC-1999; 99WO-US030311.  
PR 20-DEC-1999; 99WO-US030999.  
PR 22-DEC-1999; 99WO-US030720.  
PR 30-DEC-1999; 99WO-US031243.  
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PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000277.  
PR 11-FEB-2000; 2000WO-US003365.  
PR 18-FEB-2000; 2000WO-US004341.  
PR 22-FEB-2000; 2000WO-US004342.  
PR 24-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 01-MAR-2000; 2000WO-US005004.  
PR 02-MAR-2000; 2000WO-US005061.  
PR 02-MAR-2000; 2000WO-US005746.  
PR 10-MAR-2000; 2000WO-US005841.  
PR 15-MAR-2000; 2000WO-US006319.  
PR 20-MAR-2000; 2000WO-US006884.  
PR 21-MAR-2000; 2000WO-US007377.  
PR 30-MAR-2000; 2000WO-US007532.  
PR 17-MAY-2000; 2000WO-US013705.  
PR 22-MAY-2000; 2000WO-US014042.  
PR 30-MAY-2000; 2000WO-US014941.  
PR 02-JUN-2000; 2000WO-US015264.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US022031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023328.  
PR 08-NOV-2000; 2000WO-US030952.  
PR 10-NOV-2000; 2000WO-US030873.  
PR 01-DEC-2000; 2000WO-US032678.  
PR 20-DEC-2000; 2000US-00747259.  
PR 20-DEC-2000; 2000WO-US034956.  
PR 28-FEB-2001; 2001US-00796498.  
PR 01-MAR-2001; 2001WO-US006520.  
PR 01-MAR-2001; 2001WO-US006666.  
PR 09-MAR-2001; 2001US-00802706.  
PR 14-MAR-2001; 2001US-00808689.  
PR 22-MAR-2001; 2001US-00816744.  
PR 05-APR-2001; 2001US-00828366.  
PR 10-MAY-2001; 2001US-00854208.  
PR 18-MAY-2001; 2001US-00854280.  
PR 18-MAY-2001; 2001US-00860216.  
PR 25-MAY-2001; 2001US-00866028.  
PR 25-MAY-2001; 2001US-00866034.  
PR 25-MAY-2001; 2001WO-US017092.  
PR 01-JUN-2001; 2001US-00872035.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 05-JUN-2001; 2001US-00874503.  
PR 14-JUN-2001; 2001US-00882636.  
PR 19-JUN-2001; 2001US-00886342.  
PR 20-JUN-2001; 2001WO-US019692.  
PR 21-JUN-2001; 2001US-00887879.  
PR 22-JUN-2001; 2001WO-US020116.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 18-JUL-2001; 2001US-00908827.  
PR 06-AUG-2001; 2001US-00924419.  
PR 09-AUG-2001; 2001US-00927796.  
PR 16-AUG-2001; 2001US-00931836.  
PR 19-DEC-2001; 2001US-00028072.  
XX (GETH ) GENENTECH INC.  
XX Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;  
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;  
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;  
XX WPI: 2003-148238/14.  
XX N-PSDB; ABX89341.  
XX Two hundred and seventy five nucleic acids encoding PRO polypeptides,  
PT useful for treating pericyte-associated tumors, diabetes and various bone  
PT and/or cartilage disorders, e.g. arthritis.  
XX Claim 12; Fig 402; 659pp; English.  
XX The invention describes an isolated human PRO polypeptide. The PRO  
CC polypeptides are useful in detecting PRO polypeptides in a sample, in  
CC linking a bioactive molecule to a cell expressing a PRO polypeptide, and  
CC in modulating at least one biological activity of a cell expressing a PRO  
CC polypeptide. PRO1312 stimulates hypertrophy of neonatal heart and is thus  
CC useful for treating cardiac insufficiency disorders. PRO1154 and PRO1186  
CC stimulate adrenal cortical capillary endothelial growth, and PRO536,  
CC PRO943, PRO828, PRO826, PRO1068 or PRO535, PRO825, PRO819, PRO1126,  
CC PRO1360 and PRO1387 induce c-fos in endothelial cells, and are thus  
CC useful for treating conditions or disorders where angiogenesis would be  
CC beneficial, e.g. wound healing and antagonist of this polypeptide are  
CC useful for treating cancerous tumors. PRO812 inhibits vascular  
CC endothelial growth factor (VEGF) stimulated proliferation of endothelial  
CC cells and is thus useful for inhibiting endothelial cell growth in  
CC mammals which would be beneficial in inhibiting tumour growth. PRO826,  
CC PRO1068, PRO1184, PRO1346 and PRO1375 stimulate proliferation of  
CC stimulated T-lymphocytes and are therapeutically useful for enhancing  
CC immune response. PRO828, PRO1068 or PRO1132 enhance survival of  
CC retinal neurons cells (PRO1132 is also enhances survival/proliferation of

rod photoreceptor cells) and therefore are useful for treating retinal disorders of injuries, e.g. retinitis pigmentosa, AMD. PRO813 and PRO11066 induce proliferation of mammalian kidney mesangial cells, and therefore are useful for treating kidney disorders associated with decreased mesangial cell function such as Berger disease or other nephropathies associated with dermatitis, herpeticiformis or Crohn's disease. PRO1310, PRO844, PRO1312, PRO1192 and PRO1387 induce the proliferation and/or redifferentiation of chondrocytes in culture and are thus useful for treating sports injuries, and arthritis. This is the amino acid sequence of a novel human PRO protein

XX Sequence 123 AA;

Query Match 100.0%; Score 657; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 4.3e-62;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	1	MACRCLSLMGFLSVSQVLAQLDALLVFPQVQLSCTLSPOHVTIRDYGVSWYQQR	60
Db	1	MACRCLSLMGFLSVSQVLAQLDALLVFPQVQLSCTLSPOHVTIRDYGVSWYQQR	60
QY	61	AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNACVLTISFVQPEDDADYICSVGYG	120
Db	61	AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNACVLTISFVQPEDDADYICSVGYG	120
QY	121	FSP 123	
Db	121	FSP 123	

Search completed: June 28, 2004, 08:26:32  
Job time : 57.5135 secs

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A;Cross-references: GB:X05563; GB:Y00079; NID:G55415; PIDN:CAA9077.1; PID:G55416  
A;Note: the authors translated the codon GAG for residue 110 as Gln  
C;Superfamily: immunoglobulin V region; immunoglobulin homology  
F;20-142/Product: VpreB protein #status predicted <MAY>

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Query Match      35.5%; Score 233.5; DB 2; Length 142;
Best Local Similarity 54.7%; Pred.No.2.2e-16;
Matches 47; Conservative 9; Mismatches 29; Indels 1; Gaps 1;

33 GQAQLSCTLSQPHWITRDYGVSVYQQRAGSAPRYLLYYRSEEDHRRPADIPDRFSAKD 92
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34 GATIRLSTLNDH-NIGIYSIVYQQRGHPHPRFLRYFSDHXKQGPDIIPRFSGSKD 92
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RESULT 3  
 A28344  
 VpreB protein precursor - mouse  
 C:Species: Mus musculus (house mouse)  
 C:Date: 19-May-1989 #sequence\_revision 19-May-1989 #text\_change 21-Jul-2000  
 C:Accession: A28344  
 R:Kudo, A.; Welchers, F.  
 EMBO J. 6, 2267-2272, 1987  
 A:Title: A second gene, VpreB in the lambda-5 locus of the mouse, which appears to be se  
 A:Reference number: A91077; MUID:88029315; PMID:3117530  
 A:Accession: A28344  
 A:Molecule type: DNA  
 A:Residues: 1-142 <KUD>  
 A:Cross-references: GB:X05556; GB:Y00079; MID:g55409; PIDN:CAA29071.1; PTD:g55410  
 A:Note: the authors translated the codon GAG for residue 110 as Gln  
 C:Superfamily: immunoglobulin v region; immunoglobulin homology  
 F:20-142/Product: VpreB1 protein #status predicted <NAT>

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Query Match      34.9%; Score 229.5; DB 2; Length 142;
Best Local Similarity 53.5%;
Pred. No. 5.6e-16;
Matches 46; Conservative 9; Mismatches 30; Indels 1; Gaps 1;

QY 33 GQAQLGCTCTSPQVITRDYGVSWYQORAGSAFRLLYYRSEEDHHRPADIDPRFSAKD 92
      |||||
Db 34 GATIRLSTLNDH-NIGIYSIVYQORCHQPRFLRYSHFSKHQOEPDIPPRFSQKD 92
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RESULT 4  
PS0055  
Ig lambda chain precursor V-II region - rabbit  
C:Species: Oryctolagus cuniculus (domestic rabbit)  
C:Date: 31-Mar-1990 #sequence\_revision 31-Mar-1990 #text\_change 23-Jul-1999  
C:Accession: PS0055  
R:Hayzer, D.J.; Jaton, J.C.  
Gene 80, 185-191, 1989  
A:Title: Cloning and sequencing of two functional rabbit germ-line immunoglobulin V lambda  
A:Ref:Reference number: A91614; MUID:90006781; PMID:2507399  
A:Accession: PS0055  
A:Molecule type: DNA  
A:Residues: 1-120 <RAY>  
A:Cross-references: GB:M27840; NID:G341760; PIDN:AAA31363.1; PID:G552407  
A:Note: the authors translated the codon TTG for residue 97 as Trp  
C:Genetics: 17/1  
A:Introns: 17/1  
C:Superfamily: immunoglobulin V region; immunoglobulin homology  
C:Keywords: heterotetramer; immunoglobulin  
F:1-20/Domain: signal sequence #status predicted <SIG>  
F:21-120/Product: Ig lambda chain V-II region #status predicted <MAT>

Query Match 33.1%; Score 217.5; DB 2; Length 120;  
Best Local Similarity 41.2%; Pred. No. 7.6e-15;

	Matches	49;	Conservative	17;	Mismatches	44;	Indels	9;	Gaps	3;
Cy	5	CLSLFMGTFL-----SVSQTVLAQLDALLVPPQVAQLSCITLSPOHTTIRDYGVSWTQQR	60							
Db	3	CTPELLLLLTLCQTGSLQSPLVTQSPVSAALEGASAKLTCITLSAHT---YTIIDWYQQQ	59							
Cy	61	AGSAPRYLYVRSEEDHHRPADIPDPFSAKDEAHNAACVLTIISPVPQEDDADYYCSGY	119							
Db	60	OGSAPRYLMOLKGDSYTKTGTPDFPFGSSSGADR--YLIIPSVOADPDADYYCGADY	116							

```

RESULT 5
VpReB protein - human
S00258
C:Species: Homo sapiens (man)
C:Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 05-Nov-1999
C:Accession: S00258
R:Bauer, S.R.; Kudo, A.; Melchers, F.
EMBO J. 7, 111-116, 1988
A:Title: Structure and pre-B lymphocyte restricted expression of the VpReB gene in human
A:Reference number: S00258; MUID:88156069; PMID:3258819
A:Accession: S00258
A:Molecule type: DNA
A:Residues: 1-139 <BAU>
A:Cross-references: EMBL:M34927; NID:G340304; PIDN:AAA61292.1; PID:G340305
C:Genetics:
A:Gene: GDB:VPReB1
A:Cross-references: GDB:120493; OMIM:146770
A:Map position: 22q11.2-22q11.2

```

```

19  QTVLAQDALLFPFGQVAQLSCTLSFQHVITIRGYVSWTCQAGSAPRYLLIYRSEDDHH  78
   : : : : : : : : : : : : : : : : : : : : : : : : : : : :
20  QPVLHQFPAMWSALGTTIRLTCTLRNDH-DIGYISVYVYQORECHPFRFLIRYFSQDKS  78
   : : : : : : : : : : : : : : : : : : : : : : : : : : : :

79  READIPRFAAKDEAHNAACVLITISFQVPEDDADYCSVG  118
   : : : : : : : : : : : : : : : : : : : : : : : : : : : :
79  QGPQVPRPRTSGSKDVARNRGYLSISIQLPDEAMYICAMG  118
   : : : : : : : : : : : : : : : : : : : : : : : : : : : :

```

RESULT 6  
I57832  
VPre-B protein - human  
C:Species: Homo sapiens (man)  
C:Date: 02-Jul-1996 #sequence\_revision 02-Jul-1996 #text\_change 05-Nov-1999  
C:Accession: I57832  
R:Guelpa-Fonlupt, V.; Bossy, D.; Alzari, P.; Fumoux, F.; Fougereau, M.; Schiff, C.  
Mol. Immunol. 31, 1099-1108, 1994  
A:Ritte: The human pre-B cell receptor: structural constraints for a tentative model of  
A:Reference number: I57832; PMID:95021318; PMID:7935499  
A:Accession: I57832  
A:Status: preliminary; translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 1-145 <RES>  
A:cross-references: GB:S74019; NID:G593810; PIDN:AAB32118.1; PID:G593811  
C:Genetics:  
A:Gene: Vpre-B  
A:Introns: 16/1  
C:Superfamily: immunoglobulin V region; immunoglobulin homology

Qy	19	QTVLAQDALLVFPFGQVAQUSCTLSQPQVHTIRDYGVSVYQQRAGSAPRYLLYRSEDDHH	78
Dd	20	CPVYHQPPAMSGALGTIRLTCTLTENDH-DIGVSVYQYQQRGHPREFLYRFSQSDKS	78

DB 76 NKGSGVFRFSGNSDASAGILRISGLQLEVEADYYCGTWHSNS 122  
 RESULT 11  
 LeHUST  
 Ig lambda chain V-VI region (SUT) - human  
 C:Species: Homo sapiens (man)  
 C:Date: 30-Jun-1987 #sequence\_revision 30-Jun-1987 #text\_change 02-Sep-1997  
 C:Accession: A01988  
 R:Scotomon, A.; Kyle, R.A.; Frangione, B.  
 in Amyloidosis, Glenner, G.G.; Osserman, E.F.; Benditt, E.P.; Calkins, E.; Cohn, A.S.,  
 A>Title: Light chain variable region subgroups of monoclonal immunoglobulins in amyloidosis

A:Reference number: A01988

A:Accession: A01988

A:Molecule type: protein

A:Residues: 1-111 <SOL>

C:Genetics:

A:Gene: GDB:IGLV@

A:Cross-references: GDB:119342; OMIM:147240

A:Map position: 22q11.2-22q11.2

C:Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kappa) chain disulfide bonds. In some cases, such as IGA and IGM, the subunits associate into larger disulfide bonds.

C:Superfamily: immunoglobulin V region; immunoglobulin homology

F:1-22/Region: framework 1

F:15-93/Domain: immunoglobulin homology <IMM>

F:23-35/Region: complementarity-determining 1

F:36-50/Region: framework 2

F:51-57/Region: complementarity-determining 2

F:58-91/Region: framework 3

F:92-100/Region: complementarity-determining 3

F:101-111/Region: framework 4

F:22-91/Disulfide bonds: #status predicted

Query Match 30.7%; Score 202; DB 1; Length 111;  
Best Local Similarity 45.3%; Pred. No. 2.5e-13;  
Matches 43; Conservative 15; Mismatches 31; Indels 6; Gaps 2;

QY 21 VLALDALLVFPQVAQLSCTLSPOHVTIRDYGSWYQORAGSAPRYLLYRSEEDHRRP 80

Db 3 MLTPHVSSESPGKTVIFSCITSGG--TIAGYVQWYQORPGAPTIVP---EDTQRP 56

QY 81 ADIDRFSAKDEAHNACVLITSPVQPEDDADYYC 115

Db 57 SGVPDRFSGSIDRSSNSASLTISGLQTEDEADYYC 91

RESULT 12

A32529

Ig lambda chain precursor V region (clone pDH8) - rabbit (fragment)

C:Species: Oryctolagus cuniculus (domestic rabbit)

C:Date: 07-Jun-1990 #sequence\_revision 23-Nov-1991 #text\_change 16-Aug-1996

C:Accession: A32529

R:Hayzer, D.J.; Duvoisin, R.M.; Jaton, J.C.

Biochem. J. 245, 691-697, 1987

A:Title: cDNA clones encoding rabbit immunoglobulin lambda chains. Evidence for length v

A:Reference number: A90338; MUID:88024122; PMID:3117050

A:Accession: A32529

A:Molecule type: mRNA

A:Residues: 1-118 <HAY>

A:Cross-references: GB:M25617

A:Note: the authors translated the codon TTG for residue 37 as Phe

C:Superfamily: immunoglobulin V region; immunoglobulin homology

C:Keywords: heterotetramer; immunoglobulin

Query Match 30.58; Score 200.5; DB 2; Length 118;  
Best Local Similarity 42.3%; Pred. No. 3.8e-13;  
Matches 44; Conservative 14; Mismatches 41; Indels 5; Gaps 2;

QY 16 SVSQTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGSWYQORAGSAPRYLLYRSEE 75

Db 3 SLSPFLVLTQSPSAAALGASAKLTCTLDSAKTSL---VEVYHQKGEAPRYLMWLKDG 59

QY 76 DHRPADIPRFSAAKDEAHNACVLITSPVQPEDDADYYCSVGY 119

Db 60 SYTKGTGVPRFSGSSGADR--YLISSVQADDEADYYCGVDY 101

RESULT 13

L6HUAR

Ig lambda chain V-VI region (AR) - human (tentative sequence)

C:Species: Homo sapiens (man)

C:Date: 02-Apr-1992 #sequence\_revision 02-Apr-1982 #text\_change 31-Mar-2000

C:Accession: A01987

R:Slatten, K.; Natvig, J.B.; Husby, G.; Juul, J.

Biochem. J. 195, 561-572, 1981

A:Title: The complete amino acid sequence of a prototype immunoglobulin-lambda light-chain

A:Reference number: A01987; MUID:82091000; PMID:6797401

A:Contents: amyloid protein AR

A:Accession: A01987

A:Molecule type: protein

A:Residues: 1-112 <SLS>

A:Note: about half of the lambda chain C region is missing from this protein

C:Comment: This protein was isolated from the spleen of a patient with amyloidosis.

C:Genetics:

A:Gene: GDB:IGLV@

A:Cross-references: GDB:119342; OMIM:147240

A:Map position: 22q11.2-22q11.2

C:Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kappa) chain disulfide bonds. In some cases, such as IGA and IGM, the subunits associate into larger disulfide bonds.

C:Superfamily: immunoglobulin V region; immunoglobulin homology

C:Keywords: amyloid; heterotetramer; immunoglobulin

F:15-93/Domain: immunoglobulin homology <IMM>

F:22-91/Disulfide bonds: #status predicted

Query Match 30.3%; Score 199; DB 1; Length 112;  
Best Local Similarity 44.2%; Pred. No. 5.1e-13;  
Matches 42; Conservative 18; Mismatches 29; Indels 6; Gaps 2;

QY 21 VLALDALLVFPQVAQLSCTLSPOHVTIRDYGSWYQORAGSAPRYLLYRSEEDHRRP 80

Db 3 MLTPHVSSESPGKTVIFSCITSGG--SIADSFQWYQORPGAPTIVY----DGNRP 56

QY 81 ADIDRFSAKDEAHNACVLITSPVQPEDDADYYC 115

Db 57 SGVPDRFSGSIDSSASLTISGLKTEDEADYYC 91

RESULT 14

S04525

Ig lambda chain precursor V region - human (fragment)

C:Species: Homo sapiens (man)

C:Date: 04-Dec-1992 #sequence\_revision 04-Dec-1992 #text\_change 21-Jan-2000

C:Accession: S04525

R:Alexandre, D.; Chuchana, P.; Brockly, F.; Blancher, A.; Lefranc, G.; Lefranc, M.P.

Nucleic Acids Res. 17, 3975, 1989

A:Title: First genomic sequence of a human Ig variable lambda gene belonging to subgroup

A:Reference number: S04525; MUID:89282401; PMID:2499871

A:Accession: S04525

A:Molecule type: DNA

A:Residues: 1-117 <ALE>

A:Cross-references: EMBL:X14615; NID:g33397; PIDN:CAA32769.1; PID:g736246

C:Genetics:

A:Introns: 16/1

C:Superfamily: immunoglobulin V region; immunoglobulin homology

C:Keywords: heterotetramer; immunoglobulin

F:34-110/Domain: immunoglobulin homology <IMM>

Query Match 30.0%; Score 197; DB 2; Length 117;  
Best Local Similarity 40.3%; Pred. No. 8.6e-13;  
Matches 50; Conservative 23; Mismatches 41; Indels 10; Gaps 5;

QY 1 MACR-CLSFLLMGTFLLSVQTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQ 59

Db 1 MTCSPFLTLTLHCTGSAQSVLTQPPSVSAAPGQKVTISCGSSS--DMGNVAVSWYQ 58

QY 60 RAGSAPRYLLYRSEEDHRRPADIPRFSAAKDEAHNACVLITSPVQPEDDADYYCSVGY 119

Db 59 LPGTAPKLLIY---ENNRKPSGIPDRFSGK--SGTSATLGLTGLMPDEADYYC-LAW 111

QY 120 GFSP 123

Db 112 DTSP 115

RESULT 15

S16848

Ig lambda chain V-II region precursor - human



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GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:35 ; Search time 11.0811 Seconds  
(without alignments)  
577.979 Million cell updates/sec

Title: US-09-981-876-200  
Perfect score: 657  
Sequence: 1 MACRCLSLMGLTFLSVSQT.....PVQPEDDADYICVGVGFSP 123

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 141681 seqs, 52070155 residues

Total number of hits satisfying chosen parameters: 141681

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SwissProt\_42:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	657	100.0	123	1 VPR3_HUMAN	Q9UK13 homo sapien
2	233.5	35.5	142	1 VPR2_MOUSE	P13373 mus musculus
3	229.5	34.9	142	1 VPR1_MOUSE	P13372 mus musculus
4	215.5	32.8	145	1 VPRE_HUMAN	P06317 homo sapien
5	202	30.7	111	1 LV6C_HUMAN	P06317 homo sapien
6	199	30.3	112	1 LV6A_HUMAN	P01721 homo sapien
7	186.5	28.4	111	1 LV2F_HUMAN	P01721 homo sapien
8	185.5	28.2	111	1 LV2L_HUMAN	P80422 homo sapien
9	185	28.2	111	1 LV6D_HUMAN	P06318 homo sapien
10	185	28.2	131	1 LV6E_HUMAN	P06319 homo sapien
11	184.5	28.1	111	1 LV2G_HUMAN	P01710 homo sapien
12	183.5	27.9	130	1 LV1G_HUMAN	P06316 homo sapien
13	180.5	27.5	111	1 LV1D_HUMAN	P01702 homo sapien
14	180.5	27.5	112	1 LV2K_HUMAN	P04209 homo sapien
15	180.5	27.5	117	1 LV0A_HUMAN	P04211 homo sapien
16	179	27.2	111	1 LV2I_HUMAN	P01712 homo sapien
17	177.5	27.0	109	1 LV2E_HUMAN	P01708 homo sapien
18	175	26.6	108	1 LV3A_HUMAN	P01714 homo sapien
19	175	26.6	108	1 LV5A_HUMAN	P01719 homo sapien
20	173	26.3	109	1 LV1F_HUMAN	P04208 homo sapien
21	173	26.3	111	1 LV3B_HUMAN	P08078 homo sapien
22	172.5	26.3	110	1 LV2J_HUMAN	P01713 homo sapien
23	169.5	25.6	111	1 LV2B_HUMAN	P01705 homo sapien
24	167	25.4	106	1 LV4D_HUMAN	P01718 homo sapien
25	166	25.3	106	1 LV4B_HUMAN	P01716 homo sapien
26	166	25.3	111	1 LV4C_HUMAN	P01701 homo sapien
27	165	25.1	107	1 LV1C_HUMAN	P01717 homo sapien
28	164.5	25.0	112	1 LV1B_HUMAN	P01700 homo sapien
29	163.5	24.9	111	1 LV2H_HUMAN	P01711 homo sapien
30	163.5	24.9	112	1 LV1H_HUMAN	P06887 homo sapien
31	163	24.8	106	1 LV4H_HUMAN	P01715 homo sapien
32	162.5	24.7	109	1 KV3D_HUMAN	P01622 homo sapien
33	162.5	24.7	111	1 LV2A_HUMAN	P01704 homo sapien

34	162	24.7	109	1 LV1I_HUMAN	P06888 homo sapien
35	161.5	24.6	129	1 KV3L_HUMAN	P18135 homo sapien
36	160.5	24.4	108	1 KV3A_HUMAN	P01619 homo sapien
37	159.5	24.3	111	1 LV2C_HUMAN	P01706 homo sapien
38	158.5	24.1	111	1 LV2D_HUMAN	P01707 homo sapien
39	158	24.0	113	1 LV1_CHICK	P04210 gallus gall
40	157.5	24.0	109	1 KV3B_HUMAN	P04620 homo sapien
41	157.5	24.0	109	1 KV3G_HUMAN	P04206 homo sapien
42	155.5	23.7	129	1 KV3M_HUMAN	P18136 homo sapien
43	153	23.3	112	1 LV6B_HUMAN	P01722 homo sapien
44	151.5	23.1	115	1 KV3I_HUMAN	P04433 homo sapien
45	151	23.0	106	1 LV4E_HUMAN	P06889 homo sapien

ALIGNMENTS

RESULT 1					
VPR3_HUMAN					
ID VPR3_HUMAN	STANDARD;	PRT;	123 AA.		
AC Q9UK13;					
DT 16-OCT-2001 (Rel. 40, Created)					
DT 16-OCT-2001 (Rel. 40, Last sequence update)					
DT 15-MAR-2004 (Rel. 43, Last annotation update)					
DE Pre-B lymphocyte protein 3 precursor (VpreB3 protein) (N27C7-2).					
GN VPREB3					
OS Homo sapiens (Human)					
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;					
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.					
OX NCBI_TaxID=9606;					
RN [1]					
RP SEQUENCE FROM N.A.					
RX MEDLINE=20169186; PubMed=10702669;					
RA Rosnet O., Matei M.-G., Delatre O., Schiff C.;					
RT "VPREB3: cDNA characterization and expression in human and chromosome					
RL mapping in human and mouse.";					
RL Cytogenet. Cell Genet. 87:205-208(1999).					
RN [2]					
RP SEQUENCE FROM N.A.					
RA Shimizu N., Manosima S., Kawasaki K., Sasaki T., Hosono K.;					
RT "Molecular cloning of N27C7-2 gene.";					
RN Submitted (NOV-2000) to the ENBL/Genbank/DBJ databases.					
RN [3]					
RP SEQUENCE FROM N.A.					
RT TISSUE=Testis;					
RX MEDLINE=22388257; PubMed=12477932;					
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,					
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,					
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,					
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,					
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,					
Stapleton M., Soares M.B., Bonaldo M.P., Casavant T.L., Scheetz T.E.,					
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,					
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullany S.J.,					
Bosak S.A., McSwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,					
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,					
Villalon D.K., Murry D.M., Sodergren E.J., Lu X., Gibbs R.A.,					
Fahy J., Helton B., Kettelman M., Madan A., Rodrigues S., Sanchez A.,					
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,					
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,					
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,					
Butterfield Y.S., Krzywicki M.I., Skalska U., Smailus D.E.,					
Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;					
RT "Generation and initial analysis of more than 15,000 full-length					
human and mouse cDNA sequences.";					
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).					
CC -1- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR					
CC -1- COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS					
CC -1- TISSUE SPECIFICITY: EXPRESSED IN B cell precursors. Expressed in					
CC fetal liver, bone marrow, spleen and lymph node.					
CC -1- SIMILARITY: Belongs to the immunoglobulin superfamily.					
CC -1- SIMILARITY: Contains 1 immunoglobulin-like domain.					

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 CC -----

CC EMBL; AF163825; AAF09451.1; -;  
 CC DR EMBL; AB050772; BAB83034.1; -;  
 CC DR EMBL; BC020666; AAH20666.1; -;  
 CC DR HSSP; P01709; 2MCG.  
 CC DR Genew; HGNC:12710; VPRES3.  
 CC MIM; 605017; -;  
 CC DR InterPro; IPR007110; Ig-like.  
 CC DR InterPro; IPR003596; Ig\_V.  
 CC DR Pfam; PF00047; Ig; 1.  
 CC DR SMART; SM00406; IGV; 1.  
 CC DR PROSITE; PS50835; IG-LIKE; 1.  
 CC KW Immunoglobulin domain; Signal.  
 CC FT SIGNAL 1 20  
 CC FT CHAIN 21 123  
 CC FT DOMAIN 21 123  
 CC FT DISULFID 40 115  
 CC FT BY SIMILARITY.  
 CC SQ SEQUENCE 123 AA; 13710 MW; BF09AC5196059E85 CRC64;

Query Match 100.0%; Score 657; DB 1; Length 123;  
 Best Local Similarity 100.0%; Pred. No. 1.2e-63;  
 Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACCLSFLLMGTLFVSQTVLAQDALLVPPGVAQLSTLSPQHVITRDYGVSWYQQR 60

DB 1 MACCLSFLLMGTLFVSQTVLAQDALLVPPGVAQLSTLSPQHVITRDYGVSWYQQR 60

QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISFVQPEDDADYCYGVYG 120

DB 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISFVQPEDDADYCYGVYG 120

QY 121 FSP 123

DB 121 FSP 123

# RESULT 2

VP2 MOUSE  
 ID VP2 MOUSE STANDARD; PRT; 142 AA.  
 AC P13373;  
 DT 01-JAN-1990 (Rel. 13, Created)  
 DT 01-JAN-1990 (Rel. 13, Last sequence update)  
 DT 15-MAR-2004 (Rel. 43, Last annotation update)  
 DE Immunoglobulin omega chain precursor (VpreB2 protein).  
 GN VPRES2.  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=C57BL/6 X DBA/2J;  
 RX MEDLINE=88029315; PubMed=3117530;  
 RA Kudo A., Melchers F.;  
 RT "A second gene, VpreB in the lambda 5 locus of the mouse, which  
 RT appears to be selectively expressed in pre-B lymphocytes.";  
 RL EMO J. 6:2267-2272(1987).

CC -!- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR  
 CC COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS. THIS  
 CC COMPLEX PRESUMABLY REGULATES IG GENE REARRANGEMENTS IN THE EARLY  
 CC STEPS OF B-CELL DIFFERENTIATION  
 CC -!- TISSUE SPECIFICITY: ONLY EXPRESSED BY PRE-B-CELLS.  
 CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.  
 CC -----

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 CC or send an email to [license@isb-sib.ch](mailto:license@isb-sib.ch)).  
 CC -----

CC EMBL; X05563; CAA29077.1; -;  
 CC DR PIR; B28344; B28344.  
 CC DR HSSP; P01607; IREI.  
 CC DR MGD; MGI:98937; VpreB2.  
 CC DR InterPro; IPR007110; Ig-like.  
 CC DR InterPro; IPR003596; Ig\_V.  
 CC DR Pfam; PF00047; Ig; 1.  
 CC DR SMART; SM00406; IGV; 1.  
 CC DR PROSITE; PS50835; IG-LIKE; 1.  
 CC KW Immunoglobulin domain; Signal.  
 CC FT SIGNAL 1 19  
 CC FT CHAIN 20 142  
 CC FT DOMAIN 20 41  
 CC FT DOMAIN 42 56  
 CC FT DOMAIN 57 70  
 CC FT DOMAIN 71 81  
 CC FT DOMAIN 82 115  
 CC FT DISULFID 41 115  
 CC FT BY SIMILARITY.  
 CC SQ SEQUENCE 142 AA; 16052 MW; 7EA7128A4E63D920 CRC64;

Query Match 35.5%; Score 233.5; DB 1; Length 142;  
 Best Local Similarity 54.7%; Pred. No. 4.7e-18;  
 Matches 47; Conservative 9; Mismatches 29; Indels 1; Gaps 1;

QY 33 GQVQLSTGLPQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIPDRFSAK 92

DB 34 GATITSLTSLNDH-NICIIYIYQQRPGHPFLLRYPSHSKQGPDIIPRFSSGKD 92

QY 93 EAHNAACVLTISFVQPEDDADYCYGVG 118

DB 93 TARNLGLYSISELOPEDEAVYICAVG 118

# RESULT 3

VP1 MOUSE  
 ID VP1 MOUSE STANDARD; PRT; 142 AA.  
 AC P13372;  
 DT 01-JAN-1990 (Rel. 13, Created)  
 DT 01-JAN-1990 (Rel. 13, Last sequence update)  
 DT 15-MAR-2004 (Rel. 43, Last annotation update)  
 DE Immunoglobulin iota chain precursor (VpreB1 protein).  
 GN VPRES1.  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=C57BL/6 X DBA/2J;  
 RX MEDLINE=88029315; PubMed=3117530;  
 RA Kudo A., Melchers F.;  
 RT "A second gene, VpreB in the lambda 5 locus of the mouse, which  
 RT appears to be selectively expressed in pre-B lymphocytes.";  
 RL EMO J. 6:2267-2272(1987).

CC -!- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR  
 CC COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS. THIS  
 CC COMPLEX PRESUMABLY REGULATES IG GENE REARRANGEMENTS IN THE EARLY  
 CC STEPS OF B-CELL DIFFERENTIATION  
 CC -!- TISSUE SPECIFICITY: ONLY EXPRESSED BY PRE-B-CELLS.  
 CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.  
 CC -----

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CC -----
CC EMBL; X05556; CAA29071.1; -
CC EMBL; X05557; CAA29072.1; -
CC PIR; A28344; A28344.
CC HSP; P01607; IREI.
CC MCD; MG1.9836; VpreB1.
CC GO; GO:0005886; C:plasma membrane; IPI.
CC GO; GO:0004872; F:receptor activity; IPI.
CC GO; GO:0030097; F:hemoipoiesis; IMP.
CC GO; GO:0006955; P:immune response; IPI.
CC InterPro; IPR007110; IG-like.
CC InterPro; IPR003596; IG_v.
CC Pfam; PF00047; IG; 1.
CC SMART; SM00406; IGV; 1.
CC PROSITE; PS00835; IG LIKE; 1.
KW Immunoglobulin domain; Signal.
FT SIGNAL 1 19
FT CHAIN 20 142
FT DOMAIN 20 41
FT DOMAIN 42 56
FT DOMAIN 57 70
FT DOMAIN 71 81
FT DOMAIN 82 115
FT DISULFID 41 115
FT BY SIMILARITY.
SQ SEQUENCE 142 AA; 16125 MW; 2E18BF963A0F448C CRC64;

Query Match 34.98; Score 229.5; DB 1; Length 142;
Best Local Similarity 53.5; Pred. No. 1.3e-17;
Matches 46; Conservative 9; Mismatches 30; Indels 1; Gaps 1;

Qy 33 GQVLAQLSCTLSPQHVITRDYGVSVYQVQAGSAPRYLLYRSEEDHHRPADIPDRFSAAKD 92
Db 34 GATIRLSCTLSNDH-NIGIYSIYWYQVQPGHPFRLRYFSHDKHGQDPIDPRFSGKD 92

Qy 93 EAHNACVLITSPVOPEDADYYCSVG 118
Db 93 TTRNLGYLSISELQPEDEAVYCAVG 118

RESULT 4
VPRE_HUMAN STANDARD; PRT; 145 AA.
AC P12018;
DT 01-OCT-1989 (Rel. 12, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 15-MAR-2004 (Rel. 43, Last annotation update)
DE Immunoglobulin iota chain precursor (V(pre)B protein) (VpreB protein)
DE (CD179a antigen).
GN VPREB1 OR VPREB.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=95021318; PubMed=7935499;
RA Guelpa-Fonlupt V., Bossy D., Alzari P., Fumoux F., Fougereau M.,
RA Schiff C.;
RT "The human pre-B cell receptor: structural constraints for a tentative
RT model of the pseudo-light (psi L) chain.";
RL Mol. Immunol. 31:1099-1108(1994).
RN [2]
RP SEQUENCE FROM N.A.
RX MEDLINE=97228902; PubMed=9074928;
RA Kawasaki K., Minoshima S., Mine E., Shibuya K., Shintani A.,
RA Schmits J.L., Wang J., Shimizu N.;
RA "One-negabase sequence analysis of the human immunoglobulin lambda
RT gene locus.";
RL Genome Res. 7:250-261(1997).
RN [3]
RP SEQUENCE OF 1-139 FROM N.A.
RX MEDLINE=88196069; PubMed=3258819;

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RA Bauer S.R., Kudo A., Melchers F.;
RT "Structure and pre-B lymphocyte restricted expression of the VpreB in
RT humans and conservation of its structure in other mammalian
RT species.";
RL EMBO J. 7:111-116(1988).
CC -!- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR
CC COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS. THIS
CC COMPLEX PRESUMABLY REGULATES IG GENE REARRANGEMENTS IN THE EARLY
CC STEPS OF B-CELL DIFFERENTIATION.
CC -!- SUBUNIT: Associates non-covalently with IGLL1.
CC -!- TISSUE SPECIFICITY: ONLY EXPRESSED BY PRE-B-CELLS.
CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.
CC -!- DATABASE: NAME=PROW; NOTE=PROW 1:59-63(2000);
CC WWW="http://www.ncbi.nlm.nih.gov/prov/guide/574153212.g.htm".
CC -----
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CC -----
CC EMBL; D86992; BAA19987.1; -
CC EMBL; D88270; BAA20030.1; -
CC EMBL; S74019; AAB32118.1; -
CC EMBL; M34927; AAA61292.1; -
CC PIR; I57832; I57832.
CC PIR; S00258; S00258.
CC HSP; PF0748; ZLO1.
CC Genew; HGNC:12709; VPREB1.
CC MIM; 605141; -
CC GO; GO:0005576; C:extracellular; NAS.
CC GO; GO:0003823; F:antigen binding; NAS.
CC GO; GO:0006955; P:immune response; NAS.
CC InterPro; IPR007110; IG-like.
CC InterPro; IPR003596; IG_v.
CC Pfam; PF00047; IG; 1.
CC SMART; SM00406; IGV; 1.
CC PROSITE; PS00835; IG LIKE; 1.
KW Antigen; Signal; Immunoglobulin domain.
FT SIGNAL 1 19
FT CHAIN 20 145
FT DOMAIN 20 41
FT DOMAIN 42 56
FT DOMAIN 57 70
FT DOMAIN 71 81
FT DOMAIN 82 115
FT DISULFID 41 115
FT CONFLICT 10 10
FT SEQUENCE 145 AA; 16605 MW; 19766SB13AF64D46 CRC64;

Query Match 32.8%; Score 215.5; DB 1; Length 145;
Best Local Similarity 47.0%; Pred. No. 4.1e-16;
Matches 47; Conservative 13; Mismatches 39; Indels 1; Gaps 1;

Qy 19 QTVLAQLDALLVPGVQVLAQLSCTLSPQHVITRDYGVSVYQVQAGSAPRYLLYRSEEDH 78
Db 20 QPVLHQPAMPSALGTTIRLTCLRNH-DIGVSVYVYQVQPGHPFRLRYFSQSDKS 78

Qy 79 RPADTPDRFSAKDEAHNACVLITSPVOPEDADYYCSVG 118
Db 79 QGFQVPPRFSGSKDVARNRGYLSISELQPEDEAVYCAVG 118

RESULT 5
LV6C_HUMAN STANDARD; PRT; 111 AA.
ID LV6C_HUMAN
AC P06317;
DT 01-JAN-1988 (Rel. 06, Created)
DT 01-JAN-1988 (Rel. 06, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Ig lambda chain V-VI region SUT.

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OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
 OC Mammalia; Euthera; Primates; Catarrhini; Homidae; Homo.  
 ON NCBI\_TaxID=9606;  
 RP SEQUENCE.  
 RA Solomon A., Kyle R.A., Frangione B.;  
 RT "Light chain variable region subgroups of monoclonal immunoglobulins  
 in amyloidosis AL.";  
 RL (In) Glenner G.G., Osserman E.F., Benditt E.P., Calkins E.,  
 RL Cohen A.S., Zucker-Franklin D. (eds.); New York (1986).  
 RL Amyloidosis, pp.449-462, Plenum Press, New York (1986).  
 DR PIR; A01988; L6HUST.  
 DR PDB; 1CD0; 06-MAR-00.  
 DR InterPro; IPR007110; IG-like.  
 DR InterPro; IPR003596; IG\_v.  
 DR Pfam; PF00047; ig; 1.  
 DR SMART; SM00406; IGV; 1.  
 DR PROSITE; PS0835; IG-LIKE; 1.  
 DR Immunoglobulin V region; 3D-structure.  
 KW Immunoglobulin V region; 3D-structure.  
 FT DOMAIN 1 22  
 FT DOMAIN 23 35  
 FT DOMAIN 36 50  
 FT DOMAIN 51 57  
 FT DOMAIN 58 91  
 FT DOMAIN 92 100  
 FT DOMAIN 101 111  
 FT DISULFID 22 91  
 FT NON TER 111 111  
 SQ SEQUENCE 111 AA; 12247 MW; 0941DD547D983598 CRC64;  
 Query March 30.7%; Score 202; DB 1; Length 111;  
 Best Local Similarity 45.3%; Pred. No. 8.5e-15;  
 Matches 43; Conservative 15; Mismatches 31; Indels 6; Gaps 2;  
 QY 21 VLAQDALLVPPGQVAQLSCTLSQPHVTIRDYGSVYQQRAGSAPRYLLYRSEEDHRRP 80  
 Db 3 MLTPHVSSESPGKTVITSCRSDG--TIAGYVQVQQRPGAPTIVF---ETQRP 56  
 QY 81 ADIPDRSAKDEAHNACVLTISVPQEDDADYYC 115  
 Db 57 SGVPDRFSGSIDSSNSASLTISGLQTEDEADYYC 91  
 RESULT 6  
 LV6A\_HUMAN  
 ID LV6A\_HUMAN STANDARD; PRT; 112 AA.  
 AC P01721;  
 DT 21-JUL-1986 (Rel. 01, Created)  
 DT 21-JUL-1986 (Rel. 01, Last sequence update)  
 DT 10-OCT-2003 (Rel. 42, Last annotation update)  
 DE Ig lambda chain V-VI region AR.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
 OC Mammalia; Euthera; Primates; Catarrhini; Homidae; Homo.  
 ON NCBI\_TaxID=9606;  
 RP SEQUENCE (AMYLLOID PROTEIN AR).  
 RA Stetten K., Natvig J.B., Husby G., Juul J.;  
 RT "The complete amino acid sequence of a prototype  
 immunoglobulin-lambda light-chain-type amyloid-fibril protein AR.";  
 RL Biochem. J. 195:561-572 (1981).  
 CC CC -!- MISCELLANEOUS: ABOUT HALF OF THE LAMBDA CHAIN C REGION IS MISSING  
 CC FROM THIS PROTEIN.  
 CC CC -!- MISCELLANEOUS: THIS PROTEIN WAS ISOLATED FROM THE SPLEEN OF A  
 CC PATIENT WITH AMYLOIDOSIS.  
 CC CC -!- SIMILARITY: Contains 1 immunoglobulin-like domain.  
 DR PIR; A01987; L6HUST.  
 DR HSSP; P01709; 2MCG.  
 DR GO; GO:0005576; C:extracellular; NAS.  
 DR GO; GO:0003823; F:antigen binding; NAS.  
 DR GO; GO:0006955; P:immune response; NAS.

DR InterPro; IPR007110; IG-like.  
 DR InterPro; IPR003596; IG\_v.  
 DR Pfam; PF00047; ig; 1.  
 DR SMART; SM00406; IGV; 1.  
 DR PROSITE; PS0835; IG-LIKE; 1.  
 KW Immunoglobulin V region; Amyloid.  
 FT DOMAIN 1 107  
 FT NON TER 112 112  
 SQ SEQUENCE 112 AA; 11918 MW; 570BCD9A368EF1FE CRC64;  
 Query March 30.3%; Score 199; DB 1; Length 112;  
 Best Local Similarity 44.2%; Pred. No. 1.8e-14;  
 Matches 42; Conservative 18; Mismatches 29; Indels 6; Gaps 2;  
 QY 21 VLAQDALLVPPGQVAQLSCTLSQPHVTIRDYGSVYQQRAGSAPRYLLYRSEEDHRRP 80  
 Db 3 MLTPHVSSESPGKTVITSCRSDG--SIADSFVQVQQRPGAPTIVY---DDNCRP 56  
 QY 81 ADIPDRSAKDEAHNACVLTISVPQEDDADYYC 115  
 Db 57 SGVPDRFSGSIDSSNSASLTISGLQTEDEADYYC 91  
 RESULT 7  
 LV2P\_HUMAN  
 ID LV2P\_HUMAN STANDARD; PRT; 111 AA.  
 AC P01709;  
 DT 21-JUL-1986 (Rel. 01, Created)  
 DT 21-JUL-1986 (Rel. 01, Last sequence update)  
 DT 10-OCT-2003 (Rel. 42, Last annotation update)  
 DE Ig lambda chain V-II region MGC.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
 OC Mammalia; Euthera; Primates; Catarrhini; Homidae; Homo.  
 ON NCBI\_TaxID=9606;  
 RP SEQUENCE.  
 RA Fett J.W., Deutsch H.F.;  
 RT "Primary structure of the Mgc lambda chain.";  
 RL Biochemistry 13:4102-4114 (1974).  
 CC CC -!- MISCELLANEOUS: THE MGC-TYPE C REGION APPEARS TO BE CORRELATED WITH  
 CC A VERY UNUSUAL V-REGION SUBSTITUTION, 103-THR ABOVE FOR GLY,  
 CC SUGGESTING THAT THE V-C JOINING MECHANISM IS NOT ALWAYS RANDOM.  
 CC CC -!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE KEEN+ AND MCG+  
 CC MARKERS.  
 CC CC -!- SIMILARITY: Contains 1 immunoglobulin-like domain.  
 DR PIR; A30381; L2HUMC.  
 DR PDB; 2MCG; 15-JUL-92.  
 DR PDB; 1A8J; 17-JUN-98.  
 DR PDB; 1DCL; 15-MAY-97.

```
DR GO; GO:0005576; C:extracellular; NAS.
DR GO; GO:0003823; P:antigen binding; NAS.
DR GO; GO:0006955; P:immune response; NAS.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00406; Ig; 1.
DR PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin V region; Bence-Jones protein; 3D-structure;
KW Pyroglutamate carboxylic acid.
FT DOMAIN 1 108 IG-LIKE.
FT DISULFID 22 90 PYROGLUTAMATE CARBOXYLIC ACID.
FT STRAND 5 5 BY SIMILARITY.
FT STRAND 10 12
FT STRAND 18 23
FT TURN 26 32
FT STRAND 36 40
FT TURN 42 43
FT STRAND 50 51
FT TURN 52 54
FT STRAND 55 55
FT TURN 62 63
FT STRAND 66 68
FT STRAND 72 77
FT HELIX 82 84
FT STRAND 86 93
FT STRAND 99 101
FT STRAND 105 109
FT NON_TER 111 111
SQ SEQUENCE 111 AA; 11558 MW; 7CC1D6E2FA3377BA CRC64;

Query Match 28.4%; Score 186.5; DB 1; Length 111;
Best Local Similarity 43.9%; Pred. No. 4e-13;
Matches 43; Conservative 16; Mismatches 32; Indels 7; Gaps 3;

Qy 19 QTVLAQDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEDDHH 78
Db 1 QSALTQPPSAGSLGQSVTISCTCTSDVGQYNY-VSWYQQRAGKPKVIY----EVNK 55

Qy 79 RPADIPRFSAAKDEAHNACVLTTSPVQPEDDADYCS 116
Db 56 RPSGVPRFSGSK--SGNTASLTVSGLQAEADYCS 91

RESULT 8
LV2L HUMAN STANDARD; PRT; 111 AA.
AC P80422;
DT 01-NOV-1995 (Rel. 32, Created)
DT 01-NOV-1995 (Rel. 32, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Ig gamma lambda chain V-II region DOT.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE.
RX MEDLINE=95255298; PubMed=7737190;
RA Stoppini M., Bellotti V., Negri A., Merlini G., Garver F., Ferri G.;
RT "Characterization of the two unique human anti-flavin monooxygenase
immunoglobulins."
RL Eur. J. Biochem. 228:896-893(1995).
CC -1- SIMILARITY: Contains 1 immunoglobulin-like domain.
DR HSP; P01709; 2MCG.
DR GO; GO:0005576; C:extracellular; NAS.
DR GO; GO:0003823; P:antigen binding; NAS.
DR GO; GO:0006955; P:immune response; NAS.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00406; Ig; 1.

DR PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin V region.
FT DOMAIN 1 106 IG-LIKE.
FT DISULFID 22 90 BY SIMILARITY.
FT NON_TER 111 111
SQ SEQUENCE 111 AA; 11787 MW; F358B1EA2CD7109A CRC64;

Query Match 28.2%; Score 185.5; DB 1; Length 111;
Best Local Similarity 44.8%; Pred. No. 5.1e-13;
Matches 43; Conservative 14; Mismatches 32; Indels 7; Gaps 3;

Qy 20 TVLAQDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEDDHH 79
Db 2 SALTQPPSAGSLGQSVTISCTGLPS-VVDDNFVSWYQQTGPRAPRLIY----DDSLR 56

Qy 80 PADIPRFSAAKDEAHNACVLTTSPVQPEDDADYCS 115
Db 57 RSGVPRFSGSKSDTKAA--LTISGLQPDDEATYFC 90

RESULT 9
LV6D HUMAN STANDARD; PRT; 111 AA.
AC P06318;
DT 01-JAN-1988 (Rel. 06, Created)
DT 01-JAN-1988 (Rel. 06, Last sequence update)
DT 15-JUL-1999 (Rel. 38, Last annotation update)
DE Ig lambda chain V-VI region WLT.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE.
RX MEDLINE=86122667; PubMed=4089539;
RA Dwulet F.E., Strako K., Benson M.D.;
RT "Amino acid sequence of a lambda VI primary (AL) amyloid protein
(WLT).";
RL Scand. J. Immunol. 22:653-660(1985).
DR FIR; A01989; L6HULT.
DR HSP; P01709; 2MCG.
DR GO; GO:0005576; C:extracellular; NAS.
DR GO; GO:0003823; P:antigen binding; NAS.
DR GO; GO:0006955; P:immune response; NAS.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00406; Ig; 1.
DR PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin V region.
FT DOMAIN 1 22 FRAMEWORK-1.
FT DOMAIN 23 35 COMPLEMENTARITY-DETERMINING-1.
FT DOMAIN 36 50 FRAMEWORK-2.
FT DOMAIN 51 57 COMPLEMENTARITY-DETERMINING-2.
FT DOMAIN 58 91 FRAMEWORK-3.
FT DOMAIN 92 101 COMPLEMENTARITY-DETERMINING-3.
FT DOMAIN 102 111 FRAMEWORK-4.
FT DISULFID 22 91 BY SIMILARITY.
FT NON_TER 111 111
SQ SEQUENCE 111 AA; 11966 MW; 0C88B2FE37BCE24F CRC64;

Query Match 28.2%; Score 185; DB 1; Length 111;
Best Local Similarity 45.2%; Pred. No. 5.7e-13;
Matches 38; Conservative 16; Mismatches 24; Indels 6; Gaps 2;

Qy 32 PCVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEDDHHRADIPIRFSAAK 91
Db 14 PEKTVTISCTGSG--SIGSNVQVQYQQRGAPNTVIY----ENNRQPEVDFRFGSI 67

Qy 92 DEAHNACVLTTSPVQPEDDADYCS 115
Db 68 DSSNSGASLTISGLKTEDEADYIC 91
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RT      "Amino acid sequence of human lambda chains. 3. Tryptic peptides,
RT      chymotryptic peptides, and sequence of protein Bo.";
RL      J. Biol. Chem. 245:4488-4507(1970).
CC      -I- MISCELLANEOUS: This is a Bence-Jones protein.
CC      -I- SIMILARITY: Contains 1 immunoglobulin-like domain.
DR      PIR; A01976; L2HUBO.
DR      HSSP; P01709; 2MCG.
DR      GO; GO:0005576; C:extracellular; NAS.
DR      GO; GO:0003823; F:antigen binding; NAS.
DR      GO; GO:0006955; P:immune response; NAS.
DR      InterPro; IPR007110; Ig-like.
DR      InterPro; IPR003596; Ig_v.
DR      Pfam; PF00047; Ig; 1.
DR      SMART; SM00406; Igv; 1.
DR      PROSITE; PS50835; IG_LIKE; 1.
KW      Immunoglobulin V region; Bence-Jones protein;
KW      Pyroglutamate carboxylic acid.
FT      DOMAIN 1 106 IG-LIKE.
FT      MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
FT      DISULFID 22 90 BY SIMILARITY.
FT      NON_TER 111 111
SQ      SEQUENCE 111 AA; 11785 MW; 92F5A1BF72421BAC CRC64;

Query Match 28.1%; Score 184.5; DB 1; Length 111;
Best Local Similarity 41.8%; Pred. No. 6.5e-13;
Matches 41; Conservative 16; Mismatches 34; Indels 7; Gaps 3;

QY      19 QTVLAQLDALLVFPQVQAQLSCTLSQHVTVTRDYGVSQYQQRAGSAPRYLLYRSEEDHH 78
DB      1 QSALTQPPSASGSPGQSVTISCTGSSDVGDNKY-VSYQHPGRAPKLVI-----EVSQ 55

QY      79 RPADIPRFSAAKDEAHNACVLTISPVQPEDDADYYCS 116
DB      56 RPSGVDPFRSGSKSD--NTASLTVSGLRADADYYCS 91

RESULT 12
LVIG_HUMAN
ID      LVIG_HUMAN STANDARD; PRT; 130 AA.
AC      P06316;
DT      01-JAN-1988 (Rel. 06, Created)
DT      01-JAN-1988 (Rel. 06, Last sequence update)
DT      15-JUL-1999 (Rel. 38, Last annotation update)
DE      Ig lambda chain V-I region BL2 precursor.
OS      Homo sapiens (Human).
OC      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC      Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX      NCBI_TaxID=9606;
RN      [1]
RP      SEQUENCE FROM N.A.
RX      MEDLINE=85062823; PubMed=6095199;
RA      Tsujimoto Y., Croce C.M.;
RT      "Molecular cloning of a human immunoglobulin lambda chain variable
      sequence.";
RL      Nucleic Acids Res. 12:8407-8414(1984).
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CC      -----
DR      EMBL; X01147; CA25598.1; -
DR      PIR; A01966; L1HUBL.
DR      HSSP; P01703; 7FAB.
DR      GO; GO:0005576; C:extracellular; NAS.
DR      GO; GO:0003823; F:antigen binding; NAS.
DR      GO; GO:0006955; P:immune response; NAS.
DR      InterPro; IPR007110; Ig-like.
DR      InterPro; IPR003596; Ig_v.
DR      Pfam; PF00047; Ig; 1.

Query Match 28.2%; Score 185; DB 1; Length 131;
Best Local Similarity 42.1%; Pred. No. 6.9e-13;
Matches 40; Conservative 16; Mismatches 33; Indels 6; Gaps 2;

QY      21 VLAQLDALLVFPQVQAQLSCTLSQHVTVTRDYGVSQYQQRAGSAPRYLLYRSEEDHHP 80
DB      22 MLTQPHSVSSPGKTVTISCT--GNSGSIASNVQYQQRVSAPTVIY-----EDNQRP 75

QY      81 ADIPRFSAAKDEAHNACVLTISPVQPEDDADYYC 115
DB      76 LGVDPFRSGSIDSSNSGSLTISGLKTEADADYYC 110

RESULT 11
LV2G_HUMAN
ID      LV2G_HUMAN STANDARD; PRT; 111 AA.
AC      P01710;
DT      21-JUL-1986 (Rel. 01, Created)
DT      21-JUL-1986 (Rel. 01, Last sequence update)
DT      10-OCT-2003 (Rel. 42, Last annotation update)
DE      Ig lambda chain V-II region SO.
OS      Homo sapiens (Human).
OC      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC      Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX      NCBI_TaxID=9606;
RN      [1]
RP      SEQUENCE.
RX      MEDLINE=71103825; PubMed=5532228;
RA      Wikier M., Putnam F.W.;

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```
DR SMART, SM00406; IGV; 1.
KW PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin V region; Signal.
FT SIGNAL 1 19
FT CHAIN 20 130 IG LAMBDA CHAIN V-I REGION BL2.
FT DOMAIN 20 115 V SEGMENT.
FT DOMAIN 116 130 J SEGMENT.
FT DISULFID 41 108 BY SIMILARITY.
FT NON TER 130 130
SQ SEQUENCE 130 AA; 13564 MW; FA44BB17D3A5EBF CRC64;

Query Match 27.9%; Score 183.5; DB 1; Length 130;
Best Local Similarity 40.5%; Pred. No. 1e-12;
Matches 47; Conservative 22; Mismatches 38; Indels 9; Gaps 5;

QY 1 MACR-CLSFLLMGTFELSVSTGLVLAQLDALLVFPQVQAQLSCTLSPOHVTIRDYGVSWYQQ 59
Db 1 MTCSPLLTLLIHCSTGWSAVLTQPPSVAAAGQKVTISCSGSSNIG-NDY-VSWYQQ 58

QY 60 RAGSAAPRYLLYRSEEDHHPADIPDRFSAAKDEAHNACVLTISPVQPEDADYYC 115
Db 59 VPGTAPKLLIY----DNKPSGIPDRFSGSK--SGTSATLGTGLQTGDEADYYC 108

RESULT 13
LV1D HUMAN
ID LV1D_HUMAN STANDARD; PRT; 111 AA.
AC P01702;
DT 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Ig lambda chain V-I region NIG-64.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE.
RX MEDLINE=63186114; PubMed=6404900;
RA Kametani F., Takayasu T., Suzuki S., Shinoda T., Okuyama T.,
RT Shimizu A.;
RT "Comparative studies on the structure of the light chains of human
RT immunoglobulins. IV. Assignment of a subgroup.",
RL J. Biochem. 93:421-429(1983).
CC 1- SIMILARITY: Contains 1 immunoglobulin-like domain.
DR PIR; A01965; LIHUNG.
DR HSSP; P01703; 7FAB.
DR GO; GO:0005576; C:extracellular; NAS.
DR GO; GO:0003823; F:antigen binding; NAS.
DR GO; GO:0006955; P:immune response; NAS.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003596; IG_v.
DR Pfam; PF00047; Igv; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin V region; Pyridoxine carboxylic acid.
FT DOMAIN 1 105 IG-LIKE.
FT MOD RES 1 105 PYRROLIDONE CARBOXYLIC ACID.
FT DISULFID 22 89 BY SIMILARITY.
FT NON TER 111 111
SQ SEQUENCE 111 AA; 11454 MW; A21C6121C18A61E30 CRC64;

Query Match 27.5%; Score 180.5; DB 1; Length 111;
Best Local Similarity 41.1%; Pred. No. 1.7e-12;
Matches 44; Conservative 19; Mismatches 29; Indels 15; Gaps 4;

QY 19 QTVLAQLDALLVFPQVQAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHH 78
Db 1 QSVLTQPPSVAAAPQEQVETISCSGSSN--IGDNFVSWYQQLPGTAPKLLIY----DNKK 54

QY 79 RPADIPDRFSAAKDEAHNACVLTISPVQPEDADYYC-----SVG 118
Db 55 RPSGIPDRFSGSK--SGTSATLGTGLQTGDEADYYCCTWDSSLSVG 99
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RESULT 14
LV2K HUMAN
ID LV2K_HUMAN STANDARD; PRT; 112 AA.
AC P04209;
DT 20-MAR-1987 (Rel. 04, Created)
DT 20-MAR-1987 (Rel. 04, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Ig lambda chain V-II region NIG-84.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE.
RX MEDLINE=85204383; PubMed=3922791;
RA Tonoike H., Kametani F., Hoshi A., Shinoda T., Isobe T.;
RT "Amino acid sequence of an amyloidogenic Bence Jones protein in
RT myeloma-associated systemic amyloidosis.",
RL FEBS Lett. 185:139-141(1985).
CC 1- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN ISOLATED FROM AN
CC INDIVIDUAL WITH MYELOMA-ASSOCIATED SYSTEMIC AMYLOIDOSIS.
CC 1- SIMILARITY: Contains 1 immunoglobulin-like domain.
DR PIR; A01971; L2HUNG.
DR HSSP; P01709; 2MCG.
DR GO; GO:0005576; C:extracellular; NAS.
DR GO; GO:0003823; F:antigen binding; NAS.
DR GO; GO:0006955; P:immune response; NAS.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003596; IG_v.
DR Pfam; PF00047; Igv; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin V region; Amyloid; Bence-Jones protein.
FT DOMAIN 1 102 IG-LIKE.
FT DISULFID 22 90 BY SIMILARITY.
FT NON TER 112 112
SQ SEQUENCE 112 AA; 11581 MW; 988FEF363AE1B4F3 CRC64;

Query Match 27.5%; Score 180.5; DB 1; Length 112;
Best Local Similarity 43.9%; Pred. No. 1.8e-12;
Matches 43; Conservative 16; Mismatches 32; Indels 7; Gaps 3;

QY 19 QTVLAQLDALLVFPQVQAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHH 78
Db 1 QSALTQPPASVSGSPGQSTISCTGTSDVGGYDF-VSWYQHPGKAPKLLIY----DVNS 55

QY 79 RPADIPDRFSAAKDEAHNACVLTISPVQPEDADYYCS 116
Db 56 RPSGISNRPFGSK--SGNTASLTISGLQAEDEADYYCS 91

RESULT 15
LV0A HUMAN
ID LV0A_HUMAN STANDARD; PRT; 117 AA.
AC P04211;
DT 20-MAR-1987 (Rel. 04, Created)
DT 20-MAR-1987 (Rel. 04, Last sequence update)
DT 15-JUL-1999 (Rel. 38, Last annotation update)
DE Ig lambda chain V region 4A precursor.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=85014122; PubMed=6091030;
RA Anderson M.L.M., Szajnert M.F., Kaplan J.C., McColl L.,
RA Young B.D.;
RT "The isolation of a human Ig V lambda gene from a recombinant library
RT of chromosome 22 and estimation of its copy number.",
RL Nucleic Acids Res. 12:6647-6661(1984).
```



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OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:34 ; Search time 40.4459 Seconds

(without alignments)  
959.521 Million cell updates/sec

Title: US-09-981-876-200

Perfect score: 657  
Sequence: 1 MACRCISFLMLMGTFLSVSQT.....PVQPEDDADYICVSGYGFSP 123

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1017041 seqs, 315518202 residues

Total number of hits satisfying chosen parameters: 1017041

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

1: SPREMBL\_25.\*

2: sp\_archaea.\*

3: sp\_bacteria.\*

4: sp\_fungi.\*

5: sp\_human.\*

6: sp\_invertebrate.\*

7: sp\_mhc.\*

8: sp\_organelle.\*

9: sp\_phase.\*

10: sp\_plant.\*

11: sp\_rodent.\*

12: sp\_virus.\*

13: sp\_vertebrate.\*

14: sp\_unclassified.\*

15: sp\_rvirus.\*

16: sp\_bacteriap.\*

17: sp\_archaeap.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	430	65.4	123	11 Q61243	Q61243 mus musculus
2	235.5	35.8	230	4 Q722U3	Q722U3 homo sapien
3	199	30.3	112	4 Q96JD1	Q96JD1 homo sapien
4	199	30.3	135	4 Q9H524	Q9H524 homo sapien
5	194	29.5	112	4 Q96JD2	Q96JD2 homo sapien
6	191.5	29.1	116	4 Q96JD0	Q96JD0 homo sapien
7	190.5	29.0	236	4 Q96E61	Q96E61 homo sapien
8	187	28.5	237	4 Q9WTU6	Q9WTU6 homo sapien
9	184.5	28.1	237	4 Q9WUK4	Q9WUK4 homo sapien
10	182	27.7	240	4 Q9WUK3	Q9WUK3 homo sapien
11	178	27.1	234	4 Q9N355	Q9N355 homo sapien
12	174	26.5	235	11 Q99M11	Q99M11 mus musculus
13	171	26.0	107	4 Q9NSD6	Q9NSD6 homo sapien
14	170	25.9	233	4 Q9TBC9	Q9TBC9 homo sapien
15	170	25.9	234	4 Q722U7	Q722U7 homo sapien
16	169.5	25.8	236	4 Q9NEJ1	Q9NEJ1 homo sapien

17	168	25.6	108	4 Q96SB0	Q96SB0 homo sapien
18	167.5	25.5	109	4 Q9UL86	Q9UL86 homo sapien
19	166	25.3	100	6 Q77624	Q77624 bos taurus
20	166	25.3	110	4 Q8TE63	Q8TE63 homo sapien
21	164	25.0	233	4 Q96I69	Q96I69 homo sapien
22	164	25.0	233	4 Q9N5F4	Q9N5F4 homo sapien
23	159.5	24.3	109	4 Q9UL78	Q9UL78 homo sapien
24	158.5	24.1	105	4 Q9WVJ6	Q9WVJ6 homo sapien
25	156	23.7	81	4 Q722B8	Q722B8 homo sapien
26	154.5	23.5	132	4 Q8TBD0	Q8TBD0 homo sapien
27	154	23.4	107	4 Q9UL82	Q9UL82 homo sapien
28	151	23.0	101	4 Q8IZD8	Q8IZD8 homo sapien
29	147	22.4	248	13 Q7SVU1	Q7SVU1 xenopus lae
30	145.5	22.1	131	11 Q811C3	Q811C3 mus musculus
31	140.5	21.4	108	4 Q9UL83	Q9UL83 homo sapien
32	136.5	20.8	484	11 Q8VEA0	Q8VEA0 mus musculus
33	136	20.7	129	11 Q8VDE2	Q8VDE2 mus musculus
34	135.5	20.6	109	4 Q9UL85	Q9UL85 homo sapien
35	135.5	20.6	113	11 Q8CGS1	Q8CGS1 mus musculus
36	134	20.4	97	4 Q43234	Q43234 homo sapien
37	134	20.4	107	11 Q9ER29	Q9ER29 mus musculus
38	134	20.4	235	11 Q91W12	Q91W12 mus musculus
39	134	20.4	237	13 Q7S236	Q7S236 xenopus lae
40	133.5	20.3	93	4 Q9UL76	Q9UL76 homo sapien
41	131.5	20.0	111	11 Q811U6	Q811U6 mus musculus
42	131	19.9	235	11 Q7TMK0	Q7TMK0 mus musculus
43	131	19.9	239	4 Q8NEK0	Q8NEK0 homo sapien
44	130.5	19.9	99	11 Q9JL74	Q9JL74 mus musculus
45	130.5	19.9	108	4 Q9UL79	Q9UL79 homo sapien

#### ALIGNMENTS

RESULT 1

Q61243 PRELIMINARY; PRT; 123 AA.

AC Q61243;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)  
DE 8HS20 protein precursor (Pre-B lymphocyte gene 3).  
DE VFREB3.  
OS Mus musculus (Mouse).  
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
OX NCBI\_TaxID=10090;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC STRAIN=BALB/C;  
RX MEDLINE=93259124; PubMed=8491176;  
RA Shirasawa T., Ohnishi K., Hagiwara S., Shigemoto K., Takebe Y.,  
RA Rajewsky K., Takemori T.;  
RT "A novel gene product associated with mu chains in immature B cells.";  
RL EMBO J. 12:1827-1834(1993).  
RN [2]  
RP SEQUENCE FROM N.A.  
RC STRAIN=C57BL/6J; TISSUE=Stomach;  
RX MEDLINE=21085660; PubMed=11217851;  
RA Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,  
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,  
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,  
RA Saito T., Okazaki Y., Gojocori T., Bono H., Kasukawa T., Saito R.,  
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,  
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,  
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,  
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,  
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,  
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,  
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,  
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,  
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,  
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,

RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,  
RA Suzuki H., Toyooka K., Wang K.H., Weitz C., Whitaker C., Wilming L.,  
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohatsu S.,  
RA Hayashizaki Y.,  
RT "Functional annotation of a full-length mouse cDNA collection.";  
RL Nature 409:685-690(2001).  
DR EMBL: D13208; BAA02495.1; -.  
DR EMBL: AK008794; BAB25899.1; -.  
DR PIR: S35302; S35302.  
DR HSSP: P01709; 2MCG.  
DR MGP: MGI:98938; Vpreb3.  
DR InterPro: IPR007110; IG-like.  
DR InterPro: IPR003596; IG\_V.  
DR Pfam: PF00047; IG; 1.  
DR SMART: SM00406; IG; 1.  
DR PROSITE: PS50835; IG LIKE; 1.  
FT CHAIN 20 123 8HS20 PROTEIN.  
SQ SEQUENCE 123 AA; 13400 MW; 2A1AD371D1CEE98F CRC64;

Query Match 65.4%; Score 430; DB 11; Length 123;  
Best Local Similarity 66.1%; Pred. No. 1.2e-40;  
Matches 82; Conservative 14; Mismatches 26; Indels 2; Gaps 2;

QY 1 MAC-RCJ-SFLMGNFLSVQVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQ 59  
DB 1 MACPGCJPLLLIGTFVAVQTLTPDFAVFPQDAHLSCINSHATAGDGVSWYQQ 60  
QY 60 RAGSAPRYLLYRSEEDHRRPADIPRFSAAKDEAHNACVLITISVPQEDDADYCSV 119  
DB 61 QPGSAP-HLLYYAEHRRPADIPRFSATVDAAHNACILITISVLPEDDADYCSIAH 119  
QY 120 GFSP 123  
DB 120 TFEF 123

RESULT 2  
Q722U3 PRELIMINARY; PRT; 230 AA.  
AC Q722U3  
DT 01-OCT-2003 (TrEMBLrel. 25, Created)  
DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)  
DE Hypothetical protein.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
[1]  
SEQUENCE FROM N.A.  
MEDLINE=22389257; PubMed=12477932;  
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,  
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,  
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,  
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,  
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,  
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,  
RA Brownstein M.J., Udén T.B., Tohiyuki S., Carninci P., Prange C.,  
RA Raha S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,  
RA Besak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,  
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,  
RA Villaion D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,  
RA Fahey J., Helton E., Kettelman M., Madan A., Rodrigues S., Sanchez A.,  
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,  
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,  
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,  
RA Krzywanski M.I., Skalska U., Smillus D.E., Schnerch A., Schein J.E.,  
RA Jones S.J., Marra M.A.,  
RT "Generation and initial analysis of more than 15,000 full-length human  
and mouse cDNA sequences.";  
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).  
[2]  
SEQUENCE FROM N.A.

RA Strausberg R.;  
RL Submitted (JUL-2003) to the EMBL/GenBank/DBJ databases.  
DR EMBL: BC054893; AAH54893.1; -.  
DR Hypothetical protein.  
SQ SEQUENCE 230 AA; 24853 MW; 8BE60CC824BB886E CRC64;

Query Match 35.8%; Score 235.5; DB 4; Length 230;  
Best Local Similarity 45.5%; Pred. No. 2.4e-18;  
Matches 46; Conservative 21; Mismatches 33; Indels 1; Gaps 1;

QY 17 VSOTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEED 76  
DB 8 LSQAVLTQPSLSASPGASALCTCLR-RGFVVDYRIYVYQKSGRSPYLLRHRSDSD 66  
QY 77 HRRPADIPRFSAAKDEAHNACVLITISVPQEDDADYCSV 117  
DB 67 KOQSGVPSRFSGSKDASANAGILVIGLRSEDEADYICMV 107

RESULT 3  
Q96JDI PRELIMINARY; PRT; 112 AA.  
AC Q96JDI  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)  
DE Amyloid lambda 6 light chain variable region PIP (fragment).  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
[1]  
SEQUENCE FROM N.A.  
RP TISSUE=Bone marrow;  
RA Perfetti V., Casarini S., Colli Vignarelli M., Merlini G.;  
RT "Amyloid lambda 6 light chain variable region PIP.";  
RL Submitted (MAY-2000) to the EMBL/GenBank/DBJ databases.  
DR EMBL: AF267874; AAK56586.1; -.  
DR PIR: A30323; A30323.  
DR InterPro: IPR007110; IG-like.  
DR InterPro: IPR003596; IG\_V.  
DR Pfam: PF00047; IG; 1.  
DR SMART: SM00406; IG; 1.  
DR PROSITE: PS50835; IG LIKE; 1.  
FT NON\_TER 1 112  
FT NON\_TER 112 112  
SQ SEQUENCE 112 AA; 12047 MW; 0D3085AC2356739F CRC64;

Query Match 30.3%; Score 199; DB 4; Length 112;  
Best Local Similarity 44.2%; Pred. No. 1.3e-14;  
Matches 42; Conservative 17; Mismatches 30; Indels 6; Gaps 2;

QY 21 VLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHRRP 80  
DB 3 MLTPHVSFSGKTIITISCTRSSG--SIASNYQYQORPGSAPTIVY----EDQRP 56  
QY 81 ADIPRFSAAKDEAHNACVLITISVPQEDDADYIC 115  
DB 57 SGVDFRSGSIDSSNSGASLTISGLKTEDEADYIC 91

RESULT 4  
Q9H5Z4 PRELIMINARY; PRT; 135 AA.  
AC Q9H5Z4  
DT 01-MAR-2001 (TrEMBLrel. 16, Created)  
DT 01-MAR-2001 (TrEMBLrel. 16, Last sequence update)  
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)  
DE Hypothetical protein FLJ22755.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;



```

RN  SEQUENCE FROM N.A.
RP  TISSUE=ileal mucosa;
RA  Kawakami T., Noguchi S., Itoh T., Shigeta K., Senba T., Matsumura K.,
RA  Nakajima Y., Mizuno T., Morinaga M., Tanigami A., Fujiwara T., Ono T.,
RA  Yanada K., Fujii Y., Ozaki K., Hirao M., Ohmori Y., Ota T., Suzuki Y.,
RA  Obayashi M., Nishi T., Shibahara T., Tanaka T., Nakamura Y.,
RA  Isegai T., Sugano S.,
RA  "NEDO human cDNA sequencing project.";
RL  Submitted (AUG-2000) to the EMBL/GenBank/DBJ databases.
DR  EMBL; AK026408; BAB15473.1; -.
DR  HESP; P01607; IREI.
DR  InterPro; IPR007110; Ig-like.
DR  SMART; SMO0406; IGV; 1.
DR  Hypothetical protein.
KW  Hypothetical protein.
SQ  SEQUENCE 135 AA; 14780 MW; 652492D930F401 CRC64;

Query Match      30.3%; Score 199; DB 4; Length 135;
Best Local Similarity 45.3%; Pred. No. 1.7e-14;
Matches 34; Conservative 17; Mismatches 24; Indels 0; Gaps 0;

QY  48 TIRGYGWSVQORAGSAPRLLYYRSEEDHRRADIPDRSAKDEAHNACVLTITSPVQ 107
Db  : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
   7 SVGFWRWYQCKPGNPPRYLLYHSDNKGQGVPSRFSGSDASANAGILRLISGLQP 66

QY  108 EDDADYYCSVGYGFS 122
Db  : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
   67 EDEADYYCGTWHNS 81

RESULT 5
Q96JD2 PRELIMINARY; PRT; 112 AA.
AC Q96JD2;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Amyloid lambda 6 light chain variable region NEG (Fragment).
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Bone marrow;
RA Perfetti V., Casarini S., Colli Vignarelli M., Merlini G.;
RT "Amyloid lambda 6 light chain variable region NEG.";
RL Submitted (MAY-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL; AF267873; AAX58585.1; -.
DR InterPro; IPR007110; Ig-like.
DR SMART; SMO0406; IGV; 1.
DR PROSITE; PS50835; IG_LIKE; 1.
FT NON_TER 1
FT NON_TER 112
SQ SEQUENCE 112 AA; 11908 MW; 080B4B37E2360B06 CRC64;

Query Match      29.5%; Score 194; DB 4; Length 112;
Best Local Similarity 43.2%; Pred. No. 4.9e-14;
Matches 41; Conservative 16; Mismatches 32; Indels 6; Gaps 2;

QY  21 VLAQLDALLVFPQVAQLSCTLSPQHTIRDYGVSWYQORAGSAPRLLYYRSEEDHRRP 80
Db  3 MLTQPHSVSGPGATITISCTSGSGR--IASNSQWYQORPGSAPNIWMY----ENNQRP 56

QY  81 ADIPDRSAKDEAHNACVLTITSPVQEDDADYYC 115
Db  : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
   57 SGVDFRFGSIDSSNSASLTISGLMTEDADYYC 91

RESULT 6
Q96JD0 PRELIMINARY; PRT; 116 AA.
AC Q96JD0;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Amyloid lambda 6 light chain variable region SAR (Fragment).
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Bone marrow;
RA Perfetti V., Casarini S., Colli Vignarelli M., Merlini G.;
RT "Amyloid lambda 6 light chain variable region SAR.";
RL Submitted (MAY-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL; AF267875; AAK58587.1; -.
DR InterPro; IPR007110; Ig-like.
DR SMART; SMO0406; IGV; 1.
DR PROSITE; PS50835; IG_LIKE; 1.
FT NON_TER 1
FT NON_TER 116
SQ SEQUENCE 116 AA; 12294 MW; F7B0E9F49FAE369E CRC64;

Query Match      29.1%; Score 191.5; DB 4; Length 116;
Best Local Similarity 41.7%; Pred. No. 9.7e-14;
Matches 43; Conservative 19; Mismatches 30; Indels 11; Gaps 4;

QY  21 VLAQLDALLVFPQVAQLSCTLSPQHTIRDYGVSWYQORAGSAPRLLYYRSEEDHRRP 80
Db  3 MLTQPHSVSGPGATITISCTSGSGSIA-TNY-VQWYQLRPGSAPTIVY----EDNQRP 56

QY  81 ADIPDRSAKDEAHNACVLTITSPVQEDDADYYC----SWG 118
Db  : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
   57 SGVDFRFGSIDSSNSASLTISGLTDEADYYCQSYDSSIG 99

RESULT 7
Q95E61 PRELIMINARY; PRT; 236 AA.
AC Q95E61;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Brain;
RA Strausberg R.;
RL Submitted (AUG-2001) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC012876; AAH12876.1; -.
DR PIR; S12440; S12440.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003006; IG_MHC.
DR InterPro; IPR003596; IGV.
DR Pfam; PF00047; IGV; 2.
DR SMART; SMO0406; IGV; 1.
DR PROSITE; PS50835; IG_LIKE; 2.
DR PROSITE; PS00290; IG_MHC; 1.
DR Hypothetical protein.
KW Hypothetical protein.
SQ SEQUENCE 236 AA; 24712 MW; 7EC9FB3622FED957 CRC64;

Query Match      29.0%; Score 190.5; DB 4; Length 236;
Best Local Similarity 42.0%; Pred. No. 3e-13;
Matches 42; Conservative 21; Mismatches 30; Indels 7; Gaps 3;
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DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DE 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Brain;
RA Strausberg R.;
RL Submitted (APR-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC028090; AAH28090.1; -
DR PIR; S12441; S12441.
DR InterPro; IPR003593; Ig.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003597; Ig-cl.
DR InterPro; IPR003006; Ig_MHC.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 2.
DR SMART; SM00409; Ig; 2.
DR SMART; SM00407; IGL1; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS00835; IG_LIKE; 2.
DR PROSITE; PS00290; IG_MHC; 1.
KW Hypothetical protein.
SQ SEQUENCE 234 AA; 24792 MW; CC848CABEA4A9D63 CRC64;

Query Match 27.1%; Score 178; DB 4; Length 234;
Best Local Similarity 38.9%; Pred. No. 7.6e-12;
Matches 44; Conservative 18; Mismatches 37; Indels 14; Gaps 4;

QY 9 LLMGTEL-----SVSQTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEE 64
DB 6 LLLGLLSHCSTVTSVLTQPPSVVAPGQTARITCGN-----NIGSKSVHWYQKPGQA 61

QY 65 PRVLLYRSEEDHRRPADIPDRFSAKDEAHNACVLTISPVPQEDDADYYCSV 117
DB 62 PVLVYV-----DDSDRPSGIPERFSGS--NSGNTATLTISRVDAGDEADYYCQL 108

RESULT 12
Q99M11 ID Q99M11 PRELIMINARY; PRT; 235 AA.
AC Q99M11;
DT 01-JUN-2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DE 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RA Strausberg R.;
RL Submitted (JAN-2001) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC002129; AAH02129.1; -
DR HSSP; P01703; 7FAB.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003006; Ig_MHC.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 2.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS00835; IG_LIKE; 2.
DR PROSITE; PS00290; IG_MHC; 1.
KW Hypothetical protein.
SQ SEQUENCE 235 AA; 25403 MW; 39807BF56782A3FB CRC64;

Query Match 26.5%; Score 174; DB 11; Length 235;
Best Local Similarity 40.0%; Pred. No. 2.2e-11;
Matches 40; Conservative 16; Mismatches 38; Indels 6; Gaps 2;
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QY 16 SVSQTVLAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEE 75
DB 17 SQAQLVLTQPPSVTSLSGSLKLPCKASTGN--IGDSYVNWYQQTVMGKSPNMIY----G 70

QY 76 DHRPADIPDRFSAKDEAHNACVLTISPVPQEDDADYYC 115
DB 71 DDLRPSGVSDRFGSIDSSNSAFITIQNVQADDEADYYC 110

RESULT 13
Q9NSD6 ID Q9NSD6 PRELIMINARY; PRT; 107 AA.
AC Q9NSD6;
DT 01-OCT-2000 (TrEMBLrel. 15, Created)
DT 01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein (Fragment).
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Lymphocytes;
RA Hohmann A.;
RT "Autoimmunity.";
RL Submitted (JUL-1995) to the EMBL/GenBank/DBJ databases.
DR EMBL; L43092; AAA69746.2; -
DR HSSP; P01709; 2MCG.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS00835; IG_LIKE; 1.
DR NON_TER 1
FT NON_TER 107
SQ SEQUENCE 107 AA; 11306 MW; A2B04B37187A5F00 CRC64;

Query Match 26.0%; Score 171; DB 4; Length 107;
Best Local Similarity 40.0%; Pred. No. 1.8e-11;
Matches 38; Conservative 18; Mismatches 29; Indels 10; Gaps 3;

QY 22 LAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEEDHRRPA 81
DB 2 LTQDEVWSVALGQTVRITC---QGDSLSRSYASWYQKPGQAPVLYIGK----NNRPS 53

QY 82 DIPDRFSAKDEAHNACVLTISPVPQEDDADYYCS 116
DB 54 GIPDRFSGS--SSGNTASLTITGAQAEADYYC 86

RESULT 14
Q8TBC9 ID Q8TBC9 PRELIMINARY; PRT; 233 AA.
AC Q8TBC9;
DT 01-JUN-2002 (TrEMBLrel. 21, Created)
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=B-cell;
RA Strausberg R.;
RL Submitted (FEB-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC022823; AAH22823.1; -
DR PIR; S12442; S12442.
DR PIR; S30526; S30526.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003006; Ig_MHC.
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DR InterPro; IPR003596; Ig\_v.

DR Pfam; PF00047; Ig; 2.

DR SMART; SM00406; IGV; 1.

DR PROSITE; PS00835; IG\_LIKE; 2.

DR PROSITE; PS00290; IG\_MHC; 1.

KW Hypothetical protein.

SQ SEQUENCE 233 AA; 24867 MW; 367411BFD6F4DF92 CRC64;

Query Match 25.9%; Score 170; DB 4; Length 233;  
Best Local Similarity 40.2%; Pred.No. 6.le-11;  
Matches 39; Conservative 17; Mismatches 25; Indels 16; Gaps 4;

Qy 22 LAQDALLVFPQVQAQSCT---LSPQHVTIRYGVSWYQQRAGSAPRYLLYRSEDDHH 78

Db 23 LTQPSVSVSPGTARITCSGDALPKQY-----AYWYQKQKPGAPVLVIY----KDNE 71

Qy 79 RPADIPRFSNAKDEAHNACVLITISVPQEDDADYYC 115

Db 72 RPSGIPRFRSGS--SGTIVTLTISGVQAEDEADYYC 106

## RESULT 15

Q7ZU7

ID Q7ZU7 PRELIMINARY; PRT; 234 AA.

AC Q7ZU7;

DT 01-OCT-2003 (TrEMBLrel. 25, Created)

DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)

DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)

DE Hypothetical protein.

OS Homo sapiens (Human).

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

OX NCBI\_TaxID=9606;

RN [1]

SEQUENCE FROM N.A.

RP MEDLINE=2388257; PubMed=12477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Besak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

RA Jones S.J., Marra M.A.;

RT "Generation and initial analysis of more than 15,000 full-length human

and mouse cDNA sequences."

RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).

RN [2]

SEQUENCE FROM N.A.

RA Strausberg R.;

RL Submitted (JUL-2003) to the EMBL/GenBank/DBJ databases.

DR EMBL; BC054893; AAH54893.1; -

KW Hypothetical protein.

SQ SEQUENCE 234 AA; 25015 MW; 9A5723ABC393A06F CRC64;

Query Match 25.9%; Score 170; DB 4; Length 234;  
Best Local Similarity 36.3%; Pred.No. 6.le-11;  
Matches 41; Conservative 22; Mismatches 36; Indels 14; Gaps 4;

Qy 9 LLMGTFL-----SVSQVYLAQLDALLVFPQVQAQSCTLSFQHVITIRYGVSWYQQRAGSA 64

Db 6 LLLGLLGHCTDSVASVYLTQPPSVVAPGKTARITCGAD----NIGAKSVHWYQKTDQA 61

Qy 65 PRLLYYRSEDDHRRPADIPRFSNAKDEAHNACVLITISVPQEDDADYYCSV 117

Db 62 PVLVYH-----DDNDRPSGIPRFRSGS--NSGNTATLSISRVPGDEADYFCQV 108

Search completed: June 28, 2004, 08:27:58  
Job time : 41.4459 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: June 28, 2004, 08:28:50 ; Search time 43.7703 Seconds  
(without alignments)  
793.337 Million cell updates/sec

Title: US-09-981-876-200

Perfect score: 657  
Sequence: 1 MACRCISFLMGFLSVSQT.....PVQPEDDADYCSGVGFSF 123

Scoring table:  
Gapop 10.0 , Gapext 0.5

Searched: 1163542 seqs, 282313646 residues

Total number of hits satisfying chosen parameters: 1163542

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Published Applications AA:\*

- 1: /cgn2\_6/ptodata/2/pubpaa/US07\_PUBCOMB.pep.\*
- 2: /cgn2\_6/ptodata/2/pubpaa/PCT\_NEW\_PUB.pep.\*
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- 15: /cgn2\_6/ptodata/2/pubpaa/US10C\_PUBCOMB.pep.\*
- 16: /cgn2\_6/ptodata/2/pubpaa/US10\_NEW\_PUB.pep.\*
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- 18: /cgn2\_6/ptodata/2/pubpaa/US60\_PUBCOMB.pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query #	Score	Match	Length	DB	ID	Description
1	657	100.0	123	9	US-09-989-722-117		Sequence 117, App
2	657	100.0	123	9	US-09-989-723-117		Sequence 117, App
3	657	100.0	123	9	US-09-989-279-117		Sequence 117, App
4	657	100.0	123	9	US-09-989-727-117		Sequence 117, App
5	657	100.0	123	9	US-09-989-731-117		Sequence 117, App
6	657	100.0	123	9	US-09-989-732-117		Sequence 117, App
7	657	100.0	123	9	US-09-991-073-117		Sequence 117, App
8	657	100.0	123	9	US-09-990-442-117		Sequence 117, App
9	657	100.0	123	9	US-09-991-163-117		Sequence 117, App
10	657	100.0	123	9	US-09-993-604-117		Sequence 117, App
11	657	100.0	123	9	US-09-990-456-117		Sequence 117, App
12	657	100.0	123	9	US-09-989-721-117		Sequence 117, App
13	657	100.0	123	9	US-09-992-598-117		Sequence 117, App
14	657	100.0	123	9	US-09-981-876-200		Sequence 200, App
15	657	100.0	123	9	US-09-989-293A-117		Sequence 117, App

16	657	100.0	123	9	US-09-989-735-117		Sequence 117, App
17	657	100.0	123	9	US-09-990-444-117		Sequence 117, App
18	657	100.0	123	9	US-09-991-181-117		Sequence 117, App
19	657	100.0	123	9	US-09-989-730-117		Sequence 117, App
20	657	100.0	123	9	US-09-990-436-117		Sequence 117, App
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23	657	100.0	123	10	US-09-997-653-117		Sequence 117, App
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25	657	100.0	123	10	US-09-148-545-200		Sequence 200, App
26	657	100.0	123	10	US-09-997-428-117		Sequence 117, App
27	657	100.0	123	10	US-09-997-666-117		Sequence 117, App
28	657	100.0	123	10	US-09-990-438-117		Sequence 117, App
29	657	100.0	123	10	US-09-990-562-117		Sequence 117, App
30	657	100.0	123	10	US-09-990-711-117		Sequence 117, App
31	657	100.0	123	10	US-09-989-726-117		Sequence 117, App
32	657	100.0	123	10	US-09-988-156-117		Sequence 117, App
33	657	100.0	123	10	US-09-990-437-117		Sequence 117, App
34	657	100.0	123	10	US-09-991-157-117		Sequence 117, App
35	657	100.0	123	10	US-09-997-514-117		Sequence 117, App
36	657	100.0	123	10	US-09-997-573-117		Sequence 117, App
37	657	100.0	123	10	US-09-991-172-117		Sequence 117, App
38	657	100.0	123	10	US-09-990-726-117		Sequence 117, App
39	657	100.0	123	10	US-09-997-559-117		Sequence 117, App
40	657	100.0	123	10	US-09-997-601-117		Sequence 117, App
41	657	100.0	123	10	US-09-990-443-117		Sequence 117, App
42	657	100.0	123	10	US-09-991-854-117		Sequence 117, App
43	657	100.0	123	10	US-09-997-628-117		Sequence 117, App
44	657	100.0	123	10	US-09-997-683-117		Sequence 117, App
45	657	100.0	123	10	US-09-989-729A-117		Sequence 117, App

ALIGNMENTS

RESULT 1

US-09-989-722-117  
; Sequence 117, Application US/09989722  
; Patent No. US20020072067A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnoyers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, J. Christopher  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
; APPLICANT: Pan, James  
; APPLICANT: Paoni, Nicholas F.  
; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Williams, P. Mickey  
; APPLICANT: Wood, William I.  
; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
; FILE REFERENCE: P2730P1C63  
; CURRENT APPLICATION NUMBER: US/09/989,722  
; CURRENT FILING DATE: 2001-11-19  
; PRIOR APPLICATION NUMBER: 60/049787  
; PRIOR FILING DATE: 1997-06-16  
; PRIOR APPLICATION NUMBER: 60/062250  
; PRIOR FILING DATE: 1997-10-17

/	/	PRIOR APPLICATION NUMBER:	60/065186
/	/	PRIOR FILING DATE:	1997-11-12
/	/	PRIOR APPLICATION NUMBER:	60/065311
/	/	PRIOR FILING DATE:	1997-11-13
/	/	PRIOR APPLICATION NUMBER:	60/066770
/	/	PRIOR FILING DATE:	1997-11-24
/	/	PRIOR APPLICATION NUMBER:	60/075945
/	/	PRIOR FILING DATE:	1998-02-25
/	/	PRIOR APPLICATION NUMBER:	60/078910
/	/	PRIOR FILING DATE:	1998-03-20
/	/	PRIOR APPLICATION NUMBER:	60/083322
/	/	PRIOR FILING DATE:	1998-04-28
/	/	PRIOR APPLICATION NUMBER:	60/084600
/	/	PRIOR FILING DATE:	1998-05-07
/	/	PRIOR APPLICATION NUMBER:	60/087106
/	/	PRIOR FILING DATE:	1998-05-28
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/	/	PRIOR FILING DATE:	1998-06-02
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/	/	PRIOR FILING DATE:	1998-06-11
/	/	PRIOR APPLICATION NUMBER:	60/089105
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/	/	PRIOR APPLICATION NUMBER:	60/089440
/	/	PRIOR FILING DATE:	1998-06-16
/	/	PRIOR APPLICATION NUMBER:	60/089512

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; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
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; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091544
; PRIOR FILING DATE: 1998-07-01
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; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
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; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match          100.0%; Score 657; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 3e-64;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLSFILMGFLSVQTVLAQLDALLVFPQVAGLSCTLSPOHVTIRDYGVSWYQQR 60
Db 1 MACRCLSFILMGFLSVQTVLAQLDALLVFPQVAGLSCTLSPOHVTIRDYGVSWYQQR 60

QY 61 AGSAPRYLLYRSEEDHRRPADIPDRPSAAKDEAHNACVLTISFVQPEDDADYYCSVGYG 120
Db 61 AGSAPRYLLYRSEEDHRRPADIPDRPSAAKDEAHNACVLTISFVQPEDDADYYCSVGYG 120

QY 121 FSP 123
Db 121 FSP 123

RESULT 2
US-09-989-723-117
; Sequence 117, Application US/09989723
; Patent No. US2002072092A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730PLC62
; CURRENT APPLICATION NUMBER: US/09/989,723
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
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; PRIOR FILING DATE: 1998-06-26
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; PRIOR FILING DATE: 1998-07-01
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; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match      100.0%; Score 657; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 3e-64;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY 61 AGSAPRYLLYRSBEDHRRPADIPDRFSAKDEAHNACVLITISVPQPEDADYYCSVGYG 120
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Db 61 AGSAPRYLLYRSBEDHRRPADIPDRFSAKDEAHNACVLITISVPQPEDADYYCSVGYG 120
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QY 121 FSP 123
Db 121 FSP 123

RESULT 3
US-09-989-279-117
; Sequence 117, Application US/09989279
; Patent No. US20020072496A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tamas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C56
; CURRENT APPLICATION NUMBER: US/09/989, 279
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
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6 PRIOR FILING DATE: 1997-11-13  
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8 PRIOR FILING DATE: 1997-11-24  
9 PRIOR APPLICATION NUMBER: 60/075945  
10 PRIOR FILING DATE: 1998-02-25  
11 PRIOR APPLICATION NUMBER: 60/078910  
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13 PRIOR APPLICATION NUMBER: 60/083322  
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18 PRIOR FILING DATE: 1998-05-28  
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88 PRIOR FILING DATE: 1998-06-17  
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95 PRIOR APPLICATION NUMBER: 60/089908  
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97 PRIOR APPLICATION NUMBER: 60/089947  
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103 PRIOR APPLICATION NUMBER: 60/090246  
104 PRIOR FILING DATE: 1998-06-22  
105 PRIOR APPLICATION NUMBER: 60/090252  
106 PRIOR FILING DATE: 1998-06-22  
107 PRIOR APPLICATION NUMBER: 60/090254  
108 PRIOR FILING DATE: 1998-06-22  
109 PRIOR APPLICATION NUMBER: 60/090349  
110 PRIOR FILING DATE: 1998-06-23  
111 PRIOR APPLICATION NUMBER: 60/090355  
112 PRIOR FILING DATE: 1998-06-23  
113 PRIOR APPLICATION NUMBER: 60/090429  
114 PRIOR FILING DATE: 1998-06-24  
115 PRIOR APPLICATION NUMBER: 60/090431  
116 PRIOR FILING DATE: 1998-06-24  
117 PRIOR APPLICATION NUMBER: 60/090435  
118 PRIOR FILING DATE: 1998-06-24  
119 PRIOR APPLICATION NUMBER: 60/090444  
120 PRIOR FILING DATE: 1998-06-24  
121 PRIOR APPLICATION NUMBER: 60/090445  
122 PRIOR FILING DATE: 1998-06-24  
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132 PRIOR FILING DATE: 1998-06-24  
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135 PRIOR APPLICATION NUMBER: 60/090678  
136 PRIOR FILING DATE: 1998-06-25  
137 PRIOR APPLICATION NUMBER: 60/090690  
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139 PRIOR APPLICATION NUMBER: 60/090694  
140 PRIOR FILING DATE: 1998-06-25  
141 PRIOR APPLICATION NUMBER: 60/090695  
142 PRIOR FILING DATE: 1998-06-25  
143 PRIOR APPLICATION NUMBER: 60/090696  
144 PRIOR FILING DATE: 1998-06-25  
145 PRIOR APPLICATION NUMBER: 60/090862  
146 PRIOR FILING DATE: 1998-06-26

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; PRIOR APPLICATION NUMBER: 60/090863
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091478
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091544
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match      100.0%; Score 657; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 3e-64;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACCLSFLLMGTLFSLVSQTVAQLDALLVFPQVQVACLSTLSPOHVTIRDYGVSWYQOR 60
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Db 1 MACCLSFLLMGTLFSLVSQTVAQLDALLVFPQVQVACLSTLSPOHVTIRDYGVSWYQOR 60
   |||||

QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAAKDBAHNAACVLTISPVQPEDDADYCSGVYG 120
   |||||
Db 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAAKDBAHNAACVLTISPVQPEDDADYCSGVYG 120
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QY 121 FSP 123
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Db 121 FSP 123

RESULT 4
US-09-989-727-117
; Sequence 117, Application US/09989727
; Patent No. US2002072497A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Geritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Guiney, Austin L.
; APPLICANT: Kijavini, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paori, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1665
; CURRENT APPLICATION NUMBER: US/09/989, 727
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
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; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: 60/090863
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091478
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; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
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; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match          100.0%; Score 657; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 3e-64;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db      1  MACRCLSFLLMGTFGLSVSTVLAQLDALLVFPQVAQLSCTLSFQHWITRDYGVSWYQQR 60

QY      61  AGSAPRYLLYRSEEDHHRPADIPDRFSAAKDEAHNACVLTISPVPQEDDADYYCVSYGYG 120
Db      61  AGSAPRYLLYRSEEDHHRPADIPDRFSAAKDEAHNACVLTISPVPQEDDADYYCVSYGYG 120

QY      121  FSP 123
Db      121  FSP 123

RESULT 5
US-09-989-731-117
; Sequence 117, Application US/09989731
; Patent No. US20020103125A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
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; APPLICANT: Paoni, Nicholas P.
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; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P27301C70
; CURRENT FILING DATE: 2001-11-20
; CURRENT APPLICATION NUMBER: US/09/989, 731

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140 PRIOR FILING DATE: 1998-06-25

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; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tamas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C57
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GENERAL INFORMATION:  
APPLICANT: Askenazi, Avi J.  
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APPLICANT: Botstein, David  
APPLICANT: Desnoyers, Luc  
APPLICANT: Eaton, Dan L.  
APPLICANT: Ferrara, Napoleone  
APPLICANT: Fong, Sherman  
APPLICANT: Gerber, Hanspeter  
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APPLICANT: Wood, William I.  
APPLICANT: Zhang, Zemin  
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TITLE OF INVENTION: Acids Encodin  
FILE REFERENCE: P2730P1C15

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QY 121 FSP 123
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; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
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; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same

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; Sequence 117, Application US/09991163
; Patent No. US20020132253A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
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; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
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; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
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Query Match 100.0%; Score 657; DB 9; Length 123;
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; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
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; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tunas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730PIC25
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; CURRENT FILING DATE: 2001-11-14
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RESULT 11  
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 ; Patent No. US20020137890A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Ashkenazi, Avi J.  
 ; APPLICANT: Baker, Kevin P.  
 ; APPLICANT: Borstein, David  
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 ; APPLICANT: Watanabe, Colin K.  
 ; APPLICANT: Williams, P. Mickey  
 ; APPLICANT: Wood, William I.

APPLICANT: Zhang, Zenin  
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
FILE REFERENCE: P2730P1C22  
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Best Local Similarity 100.0%; Pred. No. 3e-64;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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DB 1 MACCLSFLLMGTLFVSQTVLAQLDALLVFPQVQLSTLSPQHVTRIDYGVSWYQQR 60

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RESULT 12
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; Patent No. US20020142961A
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
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; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
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Query Match 100.0%; Score 657; DB 9; Length 123;

Best Local Similarity 100.0%; Fred.No. 3e-64; Mismatches 0; Indels 0; Gaps 0;  
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RESULT 13

US-09-992-598-117  
; Sequence 117, Application US/09992598  
; Patent No. US20020160384A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnoyers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, J. Christopher  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
; APPLICANT: Pan, James  
; APPLICANT: Paoni, Nicholas F.  
; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.

APPLICANT: Williams, P. Mickey  
APPLICANT: Wood, William I.  
APPLICANT: Zhang, Zemin  
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
FILE REFERENCE: P2730PLC20  
CURRENT FILING DATE: 2001-11-14  
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Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 61 AGSAPRYLLYRSEEDHRPADIDPRFSAKDEAHNACVLITISVPQPEDDADYICSVGYG 120  
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## RESULT 15

US-09-989-293A-117  
; Sequence 117, Application US/09989293A  
; Patent No. US20020177164A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnoyers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, J. Christopher  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
; APPLICANT: Pan, James  
; APPLICANT: Paoni, Nicholas F.  
; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.

APPLICANT: Tumas, Daniel  
APPLICANT: Watanabe, Colin K.  
APPLICANT: Williams, P. Mickey  
APPLICANT: Wood, William I.  
APPLICANT: Zhang, Zemin  
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
TITLE OF INVENTION: Acids Encoding the Same  
FILE REFERENCE: P2730P1C66  
CURRENT APPLICATION NUMBER: US/09/989,293A  
CURRENT FILING DATE: 2001-11-20  
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; PRIOR FILING DATE: 1998-07-09

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Qy      121  FSP 123
Db      121  FSP 123

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Search completed: June 28, 2004, 08:38:21  
Job time : 44.7703 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: June 28, 2004, 08:26:40 ; Search time 197.243 Seconds  
(without alignments)  
608.663 Million cell updates/sec

Title: US-09-981-876-200

Perfect score: 657  
Sequence: 1 MACRCLSLFMGLFSLVSQT.....PVQPEDADYCSVGGRSP 123

Scoring table: EJSUM62

Gapop 10.0 , Gapext 0.5

Searched: 6019581 seqs, 976053577 residues

Total number of hits satisfying chosen parameters: 6019581

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Pending Patents\_AA\_Main.\*

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- 2: /cgn2\_6/ptodata/2/paa/US06 COMB.pcp.\*
- 3: /cgn2\_6/ptodata/2/paa/US07 COMB.pcp.\*
- 4: /cgn2\_6/ptodata/2/paa/US08 COMB.pcp.\*
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Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Length	ID	Description
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ALIGNMENTS

RESULT 1

PCT-US01-43523-402  
; Sequence 402, Application PC/TUS0143523  
; GENERAL INFORMATION:  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Heresini, Maureen  
; APPLICANT: Deforge, Laura  
; APPLICANT: Desnoyers, Luc  
; APPLICANT: Filvaroff, Ellen  
; APPLICANT: Gao, Wei-Qiang  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Smith, Victoria  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tamas, Daniel  
; APPLICANT: Watanabe, Colin K  
; APPLICANT: Wood, William  
; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC  
; TITLE OF INVENTION: ACIDS ENCODING THE SAME

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657	100.0	123	1	PCT-US02-24563-402	Sequence 402, App
657	100.0	123	20	US-09-621-011-200	Sequence 200, App
657	100.0	123	21	US-09-709-238-117	Sequence 117, App
657	100.0	123	22	US-09-791-537-2522	Sequence 2522, App
657	100.0	123	23	US-09-834-366-18461	Sequence 18461, App
657	100.0	123	24	US-09-941-992-117	Sequence 117, App
657	100.0	123	25	US-09-964-994-117	Sequence 117, App
657	100.0	123	25	US-09-981-876-200	Sequence 200, App
657	100.0	123	25	US-09-989-279-117	Sequence 117, App
657	100.0	123	25	US-09-989-293A-117	Sequence 117, App
657	100.0	123	25	US-09-989-328-117	Sequence 117, App
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657	100.0	123	25	US-09-989-722-117	Sequence 117, App
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657	100.0	123	25	US-09-990-711-117	Sequence 117, App
657	100.0	123	25	US-09-990-726-117	Sequence 117, App
657	100.0	123	25	US-09-991-073-117	Sequence 117, App
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; FILE REFERENCE: P3330R1C331
; CURRENT APPLICATION NUMBER: PCT/US01/43523
; CURRENT FILING DATE: 2002-10-02
; PRIOR APPLICATION NUMBER: 60/049911
; PRIOR FILING DATE: 1997-06-18
; PRIOR APPLICATION NUMBER: 60/056974
; PRIOR FILING DATE: 1997-08-26
; PRIOR APPLICATION NUMBER: 60/059113
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059115
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059117
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059122
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059184
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059263
; PRIOR FILING DATE: 1997-09-18
; PRIOR APPLICATION NUMBER: 60/059352
; PRIOR FILING DATE: 1997-09-19
; PRIOR APPLICATION NUMBER: 60/059588
; PRIOR FILING DATE: 1997-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 402
; LENGTH: 123
; TYPE: PRT
; ORGANISM: Homo Sapien
PCT-US01-43523-402

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Db 121 FSP 123

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; Sequence 402, Application PC/TUS0224563
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: Deforge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Geritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3330R1C331
; CURRENT APPLICATION NUMBER: PCT/US02/24563
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; PRIOR FILING DATE: 1997-09-19
; PRIOR APPLICATION NUMBER: 60/059588
; PRIOR FILING DATE: 1997-09-19
; Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 402
; LENGTH: 123
; TYPE: PRT
; ORGANISM: Homo Sapien
PCT-US02-24563-402

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; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 70 Human Secreted Proteins
; FILE REFERENCE: P2001P1
; CURRENT APPLICATION NUMBER: US/09/621,011
; CURRENT FILING DATE: 2000-07-20
; Prior application data removed - consult PALM or file wrapper
; NUMBER OF SEQ ID NOS: 280
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; SEQ ID NO 200
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-621-011-200

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QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLTISPQVEDDADYCVSVGYG 120
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; Sequence 117, Application US/09709238  
; GENERAL INFORMATION:  
; APPLICANT: Baker, Kevin  
; APPLICANT: Chen, Jian  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Gurney, Austin  
; APPLICANT: Smith, Victoria  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Wood, William I.  
; APPLICANT: Yuan, Jean  
; TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoding the Same  
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; PRIOR FILING DATE: 1998-06-25  
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; PRIOR FILING DATE: 1998-06-25  
; PRIOR APPLICATION NUMBER: US 60/090,696  
; PRIOR FILING DATE: 1998-06-25  
; PRIOR APPLICATION NUMBER: US 60/090,862  
; PRIOR FILING DATE: 1998-06-25  
; PRIOR APPLICATION NUMBER: US 60/090,863  
; PRIOR FILING DATE: 1998-06-25  
; PRIOR APPLICATION NUMBER: US 60/091,358  
; PRIOR FILING DATE: 1998-07-01  
; PRIOR APPLICATION NUMBER: US 60/091,360  
; PRIOR FILING DATE: 1998-07-01  
; PRIOR APPLICATION NUMBER: US 60/091,478  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,486  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,519  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,544  
; PRIOR FILING DATE: 1998-07-01  
; PRIOR APPLICATION NUMBER: US 60/091,626  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,628  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,633  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,646  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,673  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: US 60/091,978  
; PRIOR FILING DATE: 1998-07-07  
; PRIOR APPLICATION NUMBER: US 60/091,982  
; PRIOR FILING DATE: 1998-07-07  
; PRIOR APPLICATION NUMBER: US 60/092,182  
; PRIOR FILING DATE: 1998-07-09  
; PRIOR APPLICATION NUMBER: US 60/092,472  
; PRIOR FILING DATE: 1998-07-10  
; PRIOR APPLICATION NUMBER: US 60/093,339  
; PRIOR FILING DATE: 1998-07-20  
; PRIOR APPLICATION NUMBER: US 60/094,651  
; PRIOR FILING DATE: 1998-07-30

Query Match 100.0%; Score 657; DB 21; Length 123;  
Best Local Similarity 100.0%; Pred. No. 3.1e-65;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLSPFLMGTFLSVSQTVAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
DB 1 MACRCLSPFLMGTFLSVSQTVAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQR 60

QY 61 AGSAPRYLLYYRSEEDHRRPADIPDRFSAKDEAHNACVLTISVPQPEDDADYCVSVGYG 120  
DB 61 AGSAPRYLLYYRSEEDHRRPADIPDRFSAKDEAHNACVLTISVPQPEDDADYCVSVGYG 120  
QY 121 FSP 123  
DB 121 FSP 123

## RESULT 5

US-09-791-537-2522  
; Sequence 2522, Application US/09791537  
; GENERAL INFORMATION:  
; APPLICANT: Bionomix, Inc.  
; APPLICANT: Debe, Derek  
; APPLICANT: Danzer, Joseph  
; TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF PROTEIN FAMILIES AND FAMILY MEMBE  
; TITLE OF INVENTION: METHODS OF USE THEREOF  
; FILE REFERENCE: 261/210  
; CURRENT APPLICATION NUMBER: US/09/791,537  
; CURRENT FILING DATE: 2001-02-22  
; NUMBER OF SEQ ID NOS: 153055  
; SOFTWARE: Patent in version 3.0  
; SEQ ID NO 2522  
; LENGTH: 123  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-09-791-537-2522

Query Match 100.0%; Score 657; DB 22; Length 123;  
Best Local Similarity 100.0%; Pred. No. 3.1e-65;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLSPFLMGTFLSVSQTVAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
DB 1 MACRCLSPFLMGTFLSVSQTVAQLDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQR 60  
QY 61 AGSAPRYLLYYRSEEDHRRPADIPDRFSAKDEAHNACVLTISVPQPEDDADYCVSVGYG 120  
DB 61 AGSAPRYLLYYRSEEDHRRPADIPDRFSAKDEAHNACVLTISVPQPEDDADYCVSVGYG 120  
QY 121 FSP 123  
DB 121 FSP 123

## RESULT 6

US-09-834-366-18461  
; Sequence 18461, Application US/09834366  
; GENERAL INFORMATION:  
; APPLICANT: Bejanin, Stephane  
; APPLICANT: Tanaka, Hiroaki  
; APPLICANT: Dumas Milne Edwards, Jean Baptiste  
; APPLICANT: Jobert, Severin  
; APPLICANT: Giordano, Jean-Yves  
; TITLE OF INVENTION: ESTs and Encoded Human Proteins.  
; FILE REFERENCE: 81.US.2.REG  
; CURRENT APPLICATION NUMBER: US/09/834,366  
; CURRENT FILING DATE: 2001-04-13  
; PRIOR APPLICATION NUMBER: US 60/197,873  
; PRIOR FILING DATE: 2000-04-18  
; NUMBER OF SEQ ID NOS: 52153  
; SOFTWARE: Patent.pm  
; SEQ ID NO 18461  
; LENGTH: 123  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: SIGNAL  
; LOCATION: -20...-1  
US-09-834-366-18461

Query Match

100.0%; Score 657; DB 23; Length 123;



Best Local Similarity 100.0%; Pred. No. 3.1e-65;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLSFILMGTFLSQVTLAQDLALLVFPQVAQLCTLSPOHVTIRDYGVSNYQOR 60  
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Db 1 MACRCLSFILMGTFLSQVTLAQDLALLVFPQVAQLCTLSPOHVTIRDYGVSNYQOR 60  
|||||

QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAAKDEAHNACVLITISPVQPEDDADYYCSVGYG 120  
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Db 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAAKDEAHNACVLITISPVQPEDDADYYCSVGYG 120  
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QY 121 FSP 123  
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Db 121 FSP 123  
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## RESULT 7

US-09-941-992-117

; Sequence 117, Application US/09941992

; GENERAL INFORMATION:

; APPLICANT: Ashkenazi, Avi J.

; APPLICANT: Baker, Kevin P.

; APPLICANT: Botstein, David

; APPLICANT: Desnovers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Fong, Sherman

; APPLICANT: Gerber, Hanspeter

; APPLICANT: Gerritsen, Mary E.

; APPLICANT: Goddard, Audrey

; APPLICANT: Godowski, Paul J.

; APPLICANT: Grimaldi, J. Christopher

; APPLICANT: Gurney, Austin L.

; APPLICANT: Kljavin, Ivar J.

; APPLICANT: Napier, Mary A.

; APPLICANT: Pan, James

; APPLICANT: Paoni, Nicholas F.

; APPLICANT: Roy, Margaret Ann

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tumas, Daniel

; APPLICANT: Watanabe, Colin K.

; APPLICANT: Williams, P. Mickey

; APPLICANT: Wood, William I.

; APPLICANT: Zhang, Zemin

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

; FILE REFERENCE: P2730P1C1

; CURRENT APPLICATION NUMBER: US/09/941.992

; CURRENT FILING DATE: 2001-08-28

; PRIOR APPLICATION NUMBER: 60/049787

; PRIOR FILING DATE: 1997-06-15

; PRIOR APPLICATION NUMBER: 60/062250

; PRIOR FILING DATE: 1997-10-17

; PRIOR APPLICATION NUMBER: 60/065186

; PRIOR FILING DATE: 1997-11-12

; PRIOR APPLICATION NUMBER: 60/065311

; PRIOR FILING DATE: 1997-11-13

; PRIOR APPLICATION NUMBER: 60/066770

; PRIOR FILING DATE: 1997-11-24

; PRIOR APPLICATION NUMBER: 60/075945

; PRIOR FILING DATE: 1998-02-25

; PRIOR APPLICATION NUMBER: 60/078910

; PRIOR FILING DATE: 1998-03-20

; PRIOR APPLICATION NUMBER: 60/083322

; PRIOR FILING DATE: 1998-04-28

; PRIOR APPLICATION NUMBER: 60/084600

; PRIOR FILING DATE: 1998-05-07

; PRIOR APPLICATION NUMBER: 60/087106

; PRIOR FILING DATE: 1998-05-28

; PRIOR APPLICATION NUMBER: 60/087607

; PRIOR FILING DATE: 1998-06-02

; PRIOR APPLICATION NUMBER: 60/087609

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; PRIOR APPLICATION NUMBER: 60/088021  
; PRIOR FILING DATE: 1998-06-04  
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; PRIOR FILING DATE: 1998-06-18  
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; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09
; PRIOR APPLICATION NUMBER: 60/092472

Query Match      100.0%; Score 657; DB 24; Length 123;
Best Local Similarity 100.0%; Pred. No. 3.le-65;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 MACRCLSFLLMGTFLSVQTSLVLAQLDALLVPPGVAQLSCTLSPOHVTIRDYGVSWYQOR 60
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Db      1 MACRCLSFLLMGTFLSVQTSLVLAQLDALLVPPGVAQLSCTLSPOHVTIRDYGVSWYQOR 60
      |||

QY      61 AGSAPRYLLYRSEEDHRPADIPDRFSAAKDEAHNAACVLTIISPVQPEDDADYICSVGYG 120
      |||
Db      61 AGSAPRYLLYRSEEDHRPADIPDRFSAAKDEAHNAACVLTIISPVQPEDDADYICSVGYG 120
      |||

QY      121 RSP 123
      |||
Db      121 RSP 123
      |||

RESULT 8
US-09-964-994A-117
; Sequence 117, Application US/09964994A
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerzitsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kijavini, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: ROY, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C1
; CURRENT APPLICATION NUMBER: US/09/964,994A
; CURRENT FILING DATE: 2001-09-26
; PRIOR APPLICATION NUMBER: US/09/941,992
; PRIOR FILING DATE: 2001-08-28
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
; PRIOR APPLICATION NUMBER: 60/065311
; PRIOR FILING DATE: 1997-11-13
; PRIOR APPLICATION NUMBER: 60/066770
; PRIOR FILING DATE: 1997-11-24
; PRIOR APPLICATION NUMBER: 60/075945
; PRIOR FILING DATE: 1998-02-25
; PRIOR APPLICATION NUMBER: 60/078910
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: 60/083322
; PRIOR FILING DATE: 1998-04-28
; PRIOR APPLICATION NUMBER: 60/084600
; PRIOR FILING DATE: 1998-05-07
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 532
; SEQ ID NO 117
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; LENGTH: 123
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-964-994A-117

Query Match      100.0%; Score 657; DB 25; Length 123;
Best Local Similarity 100.0%; Pred. No. 3.1e-65;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1  MACRCLSFLLMGTFSLVSQTVLAQLDALLVFPQVQVQLSCTLSPQHVITRDYGVSWYQQR 60
Db      1  MACRCLSFLLMGTFSLVSQTVLAQLDALLVFPQVQVQLSCTLSPQHVITRDYGVSWYQQR 60

Qy      61  AGSAPRYLLYRSSEDEHRRPADIPDRFSAKDRAHNAACVLTISVPQEDDADYICVGVG 120
Db      61  AGSAPRYLLYRSSEDEHRRPADIPDRFSAKDRAHNAACVLTISVPQEDDADYICVGVG 120

Qy      121  FSP 123
Db      121  FSP 123

RESULT 9
US-09-981-876-200
; Sequence 200, Application US/09981876
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 70 Human Secreted Proteins
; FILE REFERENCE: P2001P1
; CURRENT APPLICATION NUMBER: US/09/981.876
; CURRENT FILING DATE: 2001-10-19
; PRIOR APPLICATION NUMBER: 09/148,545
; PRIOR FILING DATE: 1998-09-04
; PRIOR APPLICATION NUMBER: 60/040,162
; PRIOR FILING DATE: 1997-03-07
; PRIOR APPLICATION NUMBER: 60/040,333
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; PRIOR APPLICATION NUMBER: 60/038,621
; PRIOR FILING DATE: 1997-03-07
; PRIOR APPLICATION NUMBER: 60/040,161
; PRIOR FILING DATE: 1997-03-07
; PRIOR APPLICATION NUMBER: 60/040,626
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;; PRIOR APPLICATION NUMBER: 60/056,874  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,910  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,864  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,631  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,845  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,892  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/047,595  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/057,761  
;; PRIOR FILING DATE: 05-Sep-1997  
;; PRIOR APPLICATION NUMBER: 60/047,599  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,588  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,585  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,586  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,590  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,594  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,589  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,593  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/047,614  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/043,578  
;; PRIOR FILING DATE: 1997-04-11  
;; PRIOR APPLICATION NUMBER: 60/043,576  
;; PRIOR FILING DATE: 1997-04-11  
;; PRIOR APPLICATION NUMBER: 60/047,501  
;; PRIOR FILING DATE: 1997-05-23  
;; PRIOR APPLICATION NUMBER: 60/043,670  
;; PRIOR FILING DATE: 1997-04-11  
;; PRIOR APPLICATION NUMBER: 60/056,632  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,664  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,876  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,881  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,909  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,875  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,862  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,887  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/056,908  
;; PRIOR FILING DATE: 1997-08-22  
;; PRIOR APPLICATION NUMBER: 60/048,964  
;; PRIOR FILING DATE: 1997-06-06  
;; PRIOR APPLICATION NUMBER: 60/057,650  
;; PRIOR FILING DATE: 1997-09-05  
;; PRIOR APPLICATION NUMBER: 60/056,884  
;; PRIOR FILING DATE: 1997-08-22

;; NUMBER OF SEQ ID NOS: 280  
;; SOFTWARE: PatentIn Ver. 2.0  
;; SEQ ID NO: 200  
;; LENGTH: 123  
;; TYPE: PRT  
  
Query Match 100.0%; Score 657; DB 25; Length 123;  
Best Local Similarity 100.0%; Pred. No. 3.1e-65;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 MACRCLSFLLMGTFLLSVQTVLAQDALLVFPQVAQLSCTLSPOHVTIRDYGSVYQOR 60  
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DB 1 MACRCLSFLLMGTFLLSVQTVLAQDALLVFPQVAQLSCTLSPOHVTIRDYGSVYQOR 60  
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QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAADKDEAHNACVLTISPVPEDDADYICSVGYG 120  
|||  
DB 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAADKDEAHNACVLTISPVPEDDADYICSVGYG 120  
|||  
QY 121 FSP 123  
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DB 121 FSP 123

## RESULT 10

US-09-989-279-117  
; Sequence 117, Application US/09989279

## ; GENERAL INFORMATION:

;; APPLICANT: Ashkenazi, Avi J.  
;; APPLICANT: Baker, Kevin P.  
;; APPLICANT: Botstein, David  
;; APPLICANT: Desnoyers, Luc  
;; APPLICANT: Eaton, Dan L.  
;; APPLICANT: Ferrara, Napoleone  
;; APPLICANT: Fong, Sherman  
;; APPLICANT: Gerber, Hanspeter  
;; APPLICANT: Gerritsen, Mary E.  
;; APPLICANT: Goddard, Audrey  
;; APPLICANT: Godowski, Paul J.  
;; APPLICANT: Grimaldi, J. Christopher  
;; APPLICANT: Gurney, Austin L.  
;; APPLICANT: Kljavin, Ivar J.  
;; APPLICANT: Napier, Mary A.  
;; APPLICANT: Pan, James  
;; APPLICANT: Paoni, Nicholas F.  
;; APPLICANT: Roy, Margaret Ann  
;; APPLICANT: Stewart, Timothy A.  
;; APPLICANT: Tumas, Daniel  
;; APPLICANT: Watanabe, Colin K.  
;; APPLICANT: Williams, P. Mickey  
;; APPLICANT: Wood, William I.  
;; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
; TITLE OF INVENTION: Acids Encoding the Same  
; FILE REFERENCE: P2730P1C56  
; CURRENT APPLICATION NUMBER: US/09/989,279  
; CURRENT FILING DATE: 2001-11-19  
; PRIOR APPLICATION NUMBER: 60/049787  
; PRIOR FILING DATE: 1997-06-16  
; PRIOR APPLICATION NUMBER: 60/062250  
; PRIOR FILING DATE: 1997-10-17  
; PRIOR APPLICATION NUMBER: 60/065186  
; PRIOR FILING DATE: 1997-11-12  
; PRIOR APPLICATION NUMBER: 60/065311  
; PRIOR FILING DATE: 1997-11-13  
; PRIOR APPLICATION NUMBER: 60/066770  
; PRIOR FILING DATE: 1997-11-24  
; PRIOR APPLICATION NUMBER: 60/075945  
; PRIOR FILING DATE: 1998-02-25  
; PRIOR APPLICATION NUMBER: 60/078910  
; PRIOR FILING DATE: 1998-03-20  
; PRIOR APPLICATION NUMBER: 60/083322  
; PRIOR FILING DATE: 1998-04-28  
; PRIOR APPLICATION NUMBER: 60/084600

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; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09
; PRIOR APPLICATION NUMBER: 60/092472

Query Match          100.0%; Score 657; DB 25; Length 123;
Best Local Similarity 100.0%; Pred. No. 3.le-65;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MACRCLSFLLMGTFLSVSQTVAQLDALLVPPGQVQLSCTLSQPHVTIRDYGVSWTQQR 60
DB 1 MACRCLSFLLMGTFLSVSQTVAQLDALLVPPGQVQLSCTLSQPHVTIRDYGVSWTQQR 60

QY 61 AGSAPRYLLYRSBEDHRRPADIPDRSAKDEAHNACVLTISVPQEDDADYVCVGYG 120
DB 61 AGSAPRYLLYRSBEDHRRPADIPDRSAKDEAHNACVLTISVPQEDDADYVCVGYG 120

QY 121 FSP 123
DB 121 FSP 123

RESULT 11
US-09-989-293A-117
; Sequence 117, Application US/09989293A
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaud, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C66
; CURRENT APPLICATION NUMBER: US/09/989,293A
; CURRENT FILING DATE: 2001-11-20
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
; PRIOR APPLICATION NUMBER: 60/065311
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; PRIOR FILING DATE: 1997-11-24
; PRIOR APPLICATION NUMBER: 60/075945
; PRIOR FILING DATE: 1998-02-25
; PRIOR APPLICATION NUMBER: 60/078910
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: 60/083322
; PRIOR FILING DATE: 1998-04-28
; PRIOR APPLICATION NUMBER: 60/084600
; PRIOR FILING DATE: 1998-05-07
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; PRIOR FILING DATE: 1998-06-02
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; PRIOR FILING DATE: 1998-06-16
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; PRIOR APPLICATION NUMBER: 60/089514
; PRIOR FILING DATE: 1998-06-16
; PRIOR APPLICATION NUMBER: 60/089532
; PRIOR FILING DATE: 1998-06-17
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; PRIOR FILING DATE: 1998-06-17
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; PRIOR FILING DATE: 1998-06-17  
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; PRIOR APPLICATION NUMBER: 60/090254  
; PRIOR FILING DATE: 1998-06-22  
; PRIOR APPLICATION NUMBER: 60/090349  
; PRIOR FILING DATE: 1998-06-23  
; PRIOR APPLICATION NUMBER: 60/090355  
; PRIOR FILING DATE: 1998-06-23  
; PRIOR APPLICATION NUMBER: 60/090429  
; PRIOR FILING DATE: 1998-06-24  
; PRIOR APPLICATION NUMBER: 60/090431  
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; PRIOR FILING DATE: 1998-06-24  
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; PRIOR FILING DATE: 1998-06-24  
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; PRIOR APPLICATION NUMBER: 60/090695  
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; PRIOR FILING DATE: 1998-07-01  
; PRIOR APPLICATION NUMBER: 60/091519  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: 60/091626  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: 60/091633  
; PRIOR FILING DATE: 1998-07-02

; PRIOR APPLICATION NUMBER: 60/091978  
; PRIOR FILING DATE: 1998-07-07  
; PRIOR APPLICATION NUMBER: 60/091982  
; PRIOR FILING DATE: 1998-07-07  
; PRIOR APPLICATION NUMBER: 60/092182  
; PRIOR FILING DATE: 1998-07-09  
; PRIOR APPLICATION NUMBER: 60/092472

Query Match 100.0%; Score 657; DB 25; Length 123;  
Best Local Similarity 100.0%; Pred. No. 3.1e-65;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MACRCLSFLLMGTFLLSVSQTVLAQLDALLVFPQVLAQLSCTLSPOHVTIRDYGVSWYQOR 60  
|||||

Db 1 MACRCLSFLLMGTFLLSVSQTVLAQLDALLVFPQVLAQLSCTLSPOHVTIRDYGVSWYQOR 60  
|||||

Qy 61 AGSAPRYLLYRSEEDHHRPADIPDRFSAAKDEANACVLTITSPVQPDADADYGVSVYG 120  
|||||

Db 61 AGSAPRYLLYRSEEDHHRPADIPDRFSAAKDEANACVLTITSPVQPDADADYGVSVYG 120  
|||||

Qy 121 FSP 123  
|||

Db 121 FSP 123  
|||

## RESULT 12

US-09-989-328-117

; Sequence 117, Application US/09989328

; GENERAL INFORMATION:

; APPLICANT: Ashkenazi, Avi J.

; APPLICANT: Baker, Kevin P.

; APPLICANT: Botstein, David

; APPLICANT: Desnovers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Fong, Sherman

; APPLICANT: Gerber, Hanspeter

; APPLICANT: Gerritsen, Mary E.

; APPLICANT: Goddard, Audrey

; APPLICANT: Godowski, Paul J.

; APPLICANT: Grimaldi, J. Christopher

; APPLICANT: Gurney, Austin L.

; APPLICANT: Kljavin, Ivar J.

; APPLICANT: Napier, Mary A.

; APPLICANT: Pan, James

; APPLICANT: Paoni, Nicholas F.

; APPLICANT: Roy, Margaret Ann

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tumas, Daniel

; APPLICANT: Watanabe, Colin K.

; APPLICANT: Williams, P. Mickey

; APPLICANT: Wood, William I.

; APPLICANT: Zhang, Zemin

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

; FILE REFERENCE: P2730PIC54

; CURRENT APPLICATION NUMBER: US/09/989,328

; CURRENT FILING DATE: 2001-11-01

; PRIOR APPLICATION NUMBER: 60/049787

; PRIOR FILING DATE: 1997-06-16

; PRIOR APPLICATION NUMBER: 60/062250

; PRIOR FILING DATE: 1997-10-17

; PRIOR APPLICATION NUMBER: 60/065186

; PRIOR FILING DATE: 1997-11-12

; PRIOR APPLICATION NUMBER: 60/065311

; PRIOR FILING DATE: 1997-11-13

; PRIOR APPLICATION NUMBER: 60/066770

; PRIOR FILING DATE: 1997-11-24

; PRIOR APPLICATION NUMBER: 60/075945

; PRIOR FILING DATE: 1998-02-25

; PRIOR APPLICATION NUMBER: 60/078910

; PRIOR FILING DATE: 1998-03-20

; PRIOR APPLICATION NUMBER: 60/083322

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; PRIOR FILING DATE: 1998-04-28
; PRIOR APPLICATION NUMBER: 60/084600
; PRIOR FILING DATE: 1998-05-07
; PRIOR APPLICATION NUMBER: 60/087106
; PRIOR FILING DATE: 1998-05-28
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 532
; SEQ ID NO 117
; LENGTH: 123
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-989-328-117

Query Match          100.0%   Score 657;   DB 25;   Length 123;
Best Local Similarity 100.0%;   Pred. No. 3.le-65;
Matches 123;   Conservative 0;   Mismatches 0;   Indels 0;   Gaps 0;

QY  1  MACRCLSFLLMGTFLSVSTVLAQDLALLVFPQVQAQLSCTLSQPHVTIRDYGVSWTQQR 60
      |||||||
Db   1  MACRCLSFLLMGTFLSVSTVLAQDLALLVFPQVQAQLSCTLSQPHVTIRDYGVSWTQQR 60

QY  61  AGSAPRYLLYRSSEHHRPADIPDRFSAAXDEAHNACVLTISVPQEDDADYCVSGYG 120
      |||||||
Db   61  AGSAPRYLLYRSSEHHRPADIPDRFSAAXDEAHNACVLTISVPQEDDADYCVSGYG 120

QY  121 FSP 123
      |||
Db   121 FSP 123

RESULT 13
US-09-989-721-117
; Sequence 117, Application US/09989721
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Deenoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C55
; CURRENT APPLICATION NUMBER: US/09/989,721
; PRIOR FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
; PRIOR APPLICATION NUMBER: 60/065311
; PRIOR FILING DATE: 1997-11-13
; PRIOR APPLICATION NUMBER: 60/066770
; PRIOR FILING DATE: 1997-11-24
; PRIOR APPLICATION NUMBER: 60/075945
; PRIOR FILING DATE: 1998-02-25
; PRIOR APPLICATION NUMBER: 60/078910
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: 60/083322
; PRIOR FILING DATE: 1998-04-28
; PRIOR APPLICATION NUMBER: 60/084600
; PRIOR FILING DATE: 1998-05-07
; PRIOR APPLICATION NUMBER: 60/087106
; PRIOR FILING DATE: 1998-05-28
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/087609
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/087759
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/087827
; PRIOR FILING DATE: 1998-06-03
; PRIOR APPLICATION NUMBER: 60/088021
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088025
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088026
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088028
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; PRIOR APPLICATION NUMBER: 60/088029
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088030
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088033
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088326
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088167
; PRIOR FILING DATE: 1998-06-05
; PRIOR APPLICATION NUMBER: 60/088202
; PRIOR FILING DATE: 1998-06-05
; PRIOR APPLICATION NUMBER: 60/088212
; PRIOR FILING DATE: 1998-06-05
; PRIOR APPLICATION NUMBER: 60/088217
; PRIOR FILING DATE: 1998-06-05
; PRIOR APPLICATION NUMBER: 60/088655
; PRIOR FILING DATE: 1998-06-09
; PRIOR APPLICATION NUMBER: 60/088734
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088738
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088742
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088810
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088824
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088826
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088858
; PRIOR FILING DATE: 1998-06-11
; PRIOR APPLICATION NUMBER: 60/088861
; PRIOR FILING DATE: 1998-06-11
; PRIOR APPLICATION NUMBER: 60/088876
; PRIOR FILING DATE: 1998-06-11
; PRIOR APPLICATION NUMBER: 60/089105
; PRIOR FILING DATE: 1998-06-12
; PRIOR APPLICATION NUMBER: 60/089440
; PRIOR FILING DATE: 1998-06-16
; PRIOR APPLICATION NUMBER: 60/089512
; PRIOR FILING DATE: 1998-06-16
; PRIOR APPLICATION NUMBER: 60/089514
; PRIOR FILING DATE: 1998-06-16
; PRIOR APPLICATION NUMBER: 60/089532
; PRIOR FILING DATE: 1998-06-17
; PRIOR APPLICATION NUMBER: 60/089538
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; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09
; PRIOR APPLICATION NUMBER: 60/092472

Query Match          100.0%; Score 657; DB 25; Length 123;
Best Local Similarity 100.0%; Pred. No. 3.le-65;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1  MACRCISFLMGTFILSVSQVLAQLDALLVFPQVQVQLSCTLSPOHVTIRIDYGVSWYQQR 60
Db      1  MACRCISFLMGTFILSVSQVLAQLDALLVFPQVQVQLSCTLSPOHVTIRIDYGVSWYQQR 60

QY      61  AGAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLITISVPQEDDADYICSVGYG 120
Db      61  AGAPRYLLYRSEEDHRRPADIPDRFSAKDEAHNAACVLITISVPQEDDADYICSVGYG 120

QY      121  FSP 123
Db      121  FSP 123

RESULT 14
US-09-989-722-117
; Sequence 117, Application US/09989722
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730PlC63
; CURRENT APPLICATION NUMBER: US/09/989,722
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
; PRIOR APPLICATION NUMBER: 60/065311
; PRIOR FILING DATE: 1997-11-13
; PRIOR APPLICATION NUMBER: 60/066770

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1	PRIOR APPLICATION NUMBER: 60/089538
2	PRIOR FILING DATE: 1998-06-17
3	PRIOR APPLICATION NUMBER: 60/089598
4	PRIOR FILING DATE: 1998-06-17
5	PRIOR APPLICATION NUMBER: 60/089599
6	PRIOR FILING DATE: 1998-06-17
7	PRIOR APPLICATION NUMBER: 60/089600
8	PRIOR FILING DATE: 1998-06-17
9	PRIOR APPLICATION NUMBER: 60/089653
10	PRIOR FILING DATE: 1998-06-17
11	PRIOR APPLICATION NUMBER: 60/089801
12	PRIOR FILING DATE: 1998-06-18
13	PRIOR APPLICATION NUMBER: 60/089907
14	PRIOR FILING DATE: 1998-06-18
15	PRIOR APPLICATION NUMBER: 60/089908
16	PRIOR FILING DATE: 1998-06-18
17	PRIOR APPLICATION NUMBER: 60/089947
18	PRIOR FILING DATE: 1998-06-19
19	PRIOR APPLICATION NUMBER: 60/089948
20	PRIOR FILING DATE: 1998-06-19
21	PRIOR APPLICATION NUMBER: 60/089952
22	PRIOR FILING DATE: 1998-06-19
23	PRIOR APPLICATION NUMBER: 60/090246
24	PRIOR FILING DATE: 1998-06-22
25	PRIOR APPLICATION NUMBER: 60/090252
26	PRIOR FILING DATE: 1998-06-22
27	PRIOR APPLICATION NUMBER: 60/090254
28	PRIOR FILING DATE: 1998-06-22
29	PRIOR APPLICATION NUMBER: 60/090349
30	PRIOR FILING DATE: 1998-06-23
31	PRIOR APPLICATION NUMBER: 60/090355
32	PRIOR FILING DATE: 1998-06-23
33	PRIOR APPLICATION NUMBER: 60/090429
34	PRIOR FILING DATE: 1998-06-24
35	PRIOR APPLICATION NUMBER: 60/090431
36	PRIOR FILING DATE: 1998-06-24
37	PRIOR APPLICATION NUMBER: 60/090435
38	PRIOR FILING DATE: 1998-06-24
39	PRIOR APPLICATION NUMBER: 60/090444
40	PRIOR FILING DATE: 1998-06-24
41	PRIOR APPLICATION NUMBER: 60/090445
42	PRIOR FILING DATE: 1998-06-24
43	PRIOR APPLICATION NUMBER: 60/090472
44	PRIOR FILING DATE: 1998-06-24
45	PRIOR APPLICATION NUMBER: 60/090535
46	PRIOR FILING DATE: 1998-06-24
47	PRIOR APPLICATION NUMBER: 60/090540
48	PRIOR FILING DATE: 1998-06-24
49	PRIOR APPLICATION NUMBER: 60/090542
50	PRIOR FILING DATE: 1998-06-24
51	PRIOR APPLICATION NUMBER: 60/090557
52	PRIOR FILING DATE: 1998-06-24
53	PRIOR APPLICATION NUMBER: 60/090676
54	PRIOR FILING DATE: 1998-06-25
55	PRIOR APPLICATION NUMBER: 60/090695
56	PRIOR FILING DATE: 1998-06-25
57	PRIOR APPLICATION NUMBER: 60/090678
58	PRIOR FILING DATE: 1998-06-25
59	PRIOR APPLICATION NUMBER: 60/090690
60	PRIOR FILING DATE: 1998-06-25
61	PRIOR APPLICATION NUMBER: 60/090694
62	PRIOR FILING DATE: 1998-06-25
63	PRIOR APPLICATION NUMBER: 60/090695
64	PRIOR FILING DATE: 1998-06-25
65	PRIOR APPLICATION NUMBER: 60/090696
66	PRIOR FILING DATE: 1998-06-25
67	PRIOR APPLICATION NUMBER: 60/090862
68	PRIOR FILING DATE: 1998-06-26
69	PRIOR APPLICATION NUMBER: 60/090863
70	PRIOR FILING DATE: 1998-06-26
71	PRIOR APPLICATION NUMBER: 60/091360
72	PRIOR FILING DATE: 1998-07-01
73	PRIOR APPLICATION NUMBER: 60/091478
74	PRIOR FILING DATE: 1998-07-02
75	PRIOR APPLICATION NUMBER: 60/091544

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; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
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; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09
; PRIOR APPLICATION NUMBER: 60/092472
;
Query Match 100.0%; Score 657; DB 25; Length 123;
Best Local Similarity 100.0%; Pred. No. 3.1e-65;
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
;
QY 1 MACRCLFSLMGTFSLVSQTVLAQLDALLVFPQCVLAQLSCTLSFQHVTTIRDYGVSWYQOR 60
Db |||||
QY 61 AGSAPRYLLYRSEEDHRRPADIPDRFSAAKDEAHNACVLTISPVOPEDDADYICSVGYG 120
Db |||||
QY 121 FSP 123
Db 121 FSP 123

RESULT 15
US-09-989-723-117
; Sequence 117, Application US/09989723
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gexitsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pat, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
;
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P27301C62
; CURRENT APPLICATION NUMBER: US/09/989,723
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
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; PRIOR APPLICATION NUMBER: 60/066770
; PRIOR FILING DATE: 1997-11-24
; PRIOR APPLICATION NUMBER: 60/075945
; PRIOR FILING DATE: 1998-02-25
; PRIOR APPLICATION NUMBER: 60/078910
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: 60/083322
; PRIOR FILING DATE: 1998-04-28
; PRIOR APPLICATION NUMBER: 60/084600
; PRIOR FILING DATE: 1998-05-07
; PRIOR APPLICATION NUMBER: 60/087106
; PRIOR FILING DATE: 1998-05-28
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; PRIOR FILING DATE: 1998-06-16
; PRIOR APPLICATION NUMBER: 60/089532
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;; PRIOR FILING DATE: 1998-06-17  
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;; PRIOR APPLICATION NUMBER: 60/089598  
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;; PRIOR APPLICATION NUMBER: 60/091360  
;; PRIOR FILING DATE: 1998-07-01  
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;; PRIOR APPLICATION NUMBER: 60/091544  
;; PRIOR FILING DATE: 1998-07-01  
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;; PRIOR FILING DATE: 1998-07-02  
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;; PRIOR FILING DATE: 1998-07-02  
;; PRIOR APPLICATION NUMBER: 60/091978  
;; PRIOR FILING DATE: 1998-07-07  
;; PRIOR APPLICATION NUMBER: 60/091982  
;; PRIOR FILING DATE: 1998-07-07  
;; PRIOR APPLICATION NUMBER: 60/092182  
;; PRIOR FILING DATE: 1998-07-09  
;; PRIOR APPLICATION NUMBER: 60/092472

Query Match 100.0%; Score 657; DB 25; Length 123;  
Best Local Similarity 100.0%; Pred. No. 3.1e-65;  
Matches 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MACRCLSFLLMGTFLSVSQTVLAQLDALLVFPFGVAQLSCTLSPOHVTIRDYGVSWYQOR 60  
Db 1 MACRCLSFLLMGTFLSVSQTVLAQLDALLVFPFGVAQLSCTLSPOHVTIRDYGVSWYQOR 60  
QY 61 AGSAPRYLLYYRSEEDHRRPADIPDRFSAKDEAHNACVLTISPQPEDDADYYICSVGYG 120  
Db 61 AGSAPRYLLYYRSEEDHRRPADIPDRFSAKDEAHNACVLTISPQPEDDADYYICSVGYG 120  
QY 121 FSP 123  
Db 121 FSP 123

Search completed: June 28, 2004, 08:36:06  
Job time : 198.243 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:35 ; Search time 12.0405 seconds  
(without alignments)  
790.908 Million cell updates/sec

Title: US-09-981-876-200\_COPY\_25\_123  
Perfect score: 538  
Sequence: 1 LDALLVFPQVQAQLSCTLS.....PVPQEDADYCVSGVGGRSP 99

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 283366 seqs, 96191526 residues  
Total number of hits satisfying chosen parameters: 283366

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000  
Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : PIR 78.\*  
1: Pir1.\*  
2: Pir2.\*  
3: Pir3.\*  
4: Pir4.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Score	Match Length	DB ID	Description
1	372.5	69.2	123	B-cell protein 8HS
2	233.5	43.4	142	VpreB protein prec
3	229.5	42.7	142	VpreB protein prec
4	207.5	38.6	139	VpreB protein - hu
5	207.5	38.6	145	Vpre-B protein - h
6	200	37.2	111	Ig lambda chain V-
7	197	36.6	112	Ig lambda chain V-
8	192.5	35.8	232	Ig lambda chain pr
9	191.5	35.6	243	Ig lambda chain -
10	189.5	35.2	120	Ig lambda chain pr
11	185.5	34.5	99	Ig lambda chain -
12	185.5	34.5	111	Ig lambda chain V-
13	185	34.4	111	Ig lambda chain V-
14	184.5	34.3	136	Ig lambda chain pr
15	184.5	34.3	120	Ig lambda chain V-
16	183.5	34.1	118	Ig lambda chain pr
17	183	34.0	131	Ig lambda chain pr
18	182.5	33.9	133	Ig lambda chain (D
19	182.5	33.9	216	Ig lambda chain -
20	181.5	33.7	99	Ig lambda chain -
21	180.5	33.6	98	Ig lambda chain V-
22	180.5	33.6	111	Ig lambda chain pr
23	180.5	33.6	117	Ig lambda chain pr
24	180	33.5	107	Ig lambda chain -
25	178.5	33.2	235	Ig lambda chain -
26	176.5	32.8	94	Ig lambda chain V
27	176.5	32.8	111	Ig lambda chain V
28	176.5	32.8	112	Ig lambda chain V
29	176	32.7	108	Ig lambda chain -

30 175.5 32.6 99 2 S36051  
31 175.5 32.6 106 2 S40091  
32 175.5 32.6 112 1 L2HUNG  
33 175.5 32.6 234 2 A39956  
34 175 32.5 111 1 L2HUNW  
35 174.5 32.4 99 2 S36056  
36 173.5 32.2 111 2 S19673  
37 173.5 32.2 132 2 A55410  
38 173 32.2 112 2 D44151  
39 173 32.2 113 2 A29700  
40 172.5 32.1 99 2 S36052  
41 172.5 32.1 99 2 S36053  
42 172.5 32.1 109 1 L2HUBR  
43 172.5 32.1 112 2 S51148  
44 172.5 32.1 118 2 S12627  
45 171.5 31.9 110 2 S57412

ALIGNMENTS

RESULT 1

B-cell protein 8HS-20 precursor - mouse  
C:Species: Mus musculus (house mouse)  
C>Date: 31-Dec-1993 #sequence\_revision 02-Jun-1994 #text\_change 20-Jun-2000  
C:Accession: S35302  
R:Shitasawa, T.; Ohnishi, K.; Hagiwara, S.; Shigemoto, K.; Takebe, Y.; Rajewsky, K.; Ta  
EMBO J. 12, 1827-1834, 1993  
A>Title: A novel gene product associated with mu chains in immature B cells.  
A:Reference number: S35302; MUID:93259124; PMID:8491176  
A:Accession: S35302  
A:Molecule type: DNA  
A:Residues: 1-123 <SHI>  
A:Cross-references: EMBL: D13208; NID: G286064; PIDN: BAA02495.1; PID: G286065  
C:Genetics:  
A:Gene: 8HS-20  
A:Introns: 18/1  
C:Superfamily: immunoglobulin V region; immunoglobulin homology  
C:Keywords: B-cell  
F:1-19/Domain: signal sequence #status predicted <SIG>  
F:20-123/Product: B-cell protein 8HS-20 #status predicted <MAT>

Query Match 69.2%; Score 372.5; DB 2; Length 123;

Best Local Similarity 69.4%; Pred. No. 2.3e-31; Mismatches 18; Indels 1; Gaps 1;

2 DALLVFPQVQAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIPD 61

27 DAFSVFPGQDAHLSCITNSQHATAGDIGVSWYQQQPGSAP-HLLYYAAEEHVRPADIPD 85

62 RFSAAKDEAHNAACVLITISVPQEDDADYCVSGVGGRSP 99

86 RFSATVDAAHNAACILITISVLPEDDADYCVSIAHTFEP 123

RESULT 2

B28344

VpreB protein precursor - mouse

C:Species: Mus musculus (house mouse)

C>Date: 19-May-1989 #sequence\_revision 19-May-1989 #text\_change 05-Nov-1999

C:Accession: B28344

R:Kudo, A.; Welchers, F.

EMBO J. 6, 2267-2275, 1987

A>Title: A second gene, VpreB in the lambda-5 locus of the mouse, which appears to be s

A:Reference number: A91077; MUID:88029315; PMID:3117530

A:Accession: B28344

A:Molecule type: DNA

A:Residues: 1-142 <KUD>

A:Cross-references: GB:X05563; GB:Y00079; NID:955415; PIDN:CAA29077.1; PID:955416

A>Note: the authors translated the codon GAG for residue 110 as Gln

C:Superfamily: immunoglobulin V region; immunoglobulin homology

F:20-142/Product: VpreB protein #status predicted <MAT>

```
Query Match      43.4%; Score 233.5; DB 2; Length 142;
Best Local Similarity 54.7%; Pred. No. 5.8e-17;
Matches 47; Conservative 9; Mismatches 29; Indels 1; Gaps 1;

QY 9 GQVQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68
Db 34 GATIRLCTLSNDH-NIGIYIYVYQQRPGHPFRFLRYFSHSDKHQGPDPFRFSGSKD 92

QY 69 EAHNACVLTISPVQPEDDADYVCSVG 94
Db 93 TARNLGYLSISELQPEDEAVYCAVG 118

RESULT 3
A28344
VpreB protein precursor - mouse
C:Species: Mus musculus (house mouse)
C:Date: 19-May-1989 #sequence_revision 19-May-1989 #text_change 21-Jul-2000
C:Accession: A28344
R:Kudo, A.; Melchers, F.
EMBO J. 6, 2267-2272, 1987
A:Title: A second gene, VpreB in the lambda-5 locus of the mouse, which appears to be se
A:Reference number: A91077; MUID:89029315; PMID:31117530
A:Accession: A28344
A:Molecule type: DNA
A:Residues: 1-142 <KUD>
A:Cross-references: GB:X05556; GB:Y00079; NID:G55409; PIDN:CAA29071.1; PID:G55410
A:Note: the authors translated the codon GAG for residue 110 as Gln
C:Superfamily: immunoglobulin V region; immunoglobulin homology
F:20-142/Product: VpreB1 protein #status predicted <MAR>

Query Match      42.7%; Score 229.5; DB 2; Length 142;
Best Local Similarity 53.5%; Pred. No. 1.5e-16;
Matches 46; Conservative 9; Mismatches 30; Indels 1; Gaps 1;

QY 9 GQVQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68
Db 34 GATIRLCTLSNDH-NIGIYIYVYQQRPGHPFRFLRYFSHSDKHQGPDPFRFSGSKD 92

QY 69 EAHNACVLTISPVQPEDDADYVCSVG 94
Db 93 TTRNLGYLSISELQPEDEAVYCAVG 118

RESULT 4
S00258
VpreB protein - human
C:Species: Homo sapiens (man)
C:Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 05-Nov-1999
C:Accession: S00258
R:Bauer, S.R.; Kudo, A.; Melchers, F.
EMBO J. 7, 111-116, 1988
A:Title: Structure and pre-B lymphocyte restricted expression of the VpreB gene in human
A:Reference number: S00258; MUID:88196069; PMID:3258819
A:Accession: S00258
A:Molecule type: DNA
A:Residues: 1-139 <BAU>
A:Cross-references: EMBL:M34927; NID:G340304; PIDN:AAA61292.1; PID:G340305
C:Genetics:
A:Gene: GDB:VPREB1
A:Cross-references: GDB:120493; OMIM:146770
A:Map position: 22q11.2-22q11.2
A:Introns: 16/1
C:Superfamily: immunoglobulin V region; immunoglobulin homology

Query Match      38.6%; Score 207.5; DB 2; Length 139;
Best Local Similarity 48.8%; Pred. No. 2.7e-14;
Matches 42; Conservative 12; Mismatches 31; Indels 1; Gaps 1;

QY 9 GQVQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68
Db 34 GTTIRLCTLRNDH-DIGVSVYVYQQRPGHPFRFLRYFSQSDKSGQGPVPPRFSGSKD 92
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QY 69 EAHNACVLTISPVQPEDDADYVCSVG 94
Db 93 VARNRGYLSISELQPEDEAVYCAVG 118

RESULT 5
I57832
Vpre-B protein - human
C:Species: Homo sapiens (man)
C:Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 05-Nov-1999
C:Accession: I57832
R:Guelpa-Ponlupt, V.; Bossy, D.; Alzari, P.; Fumoux, F.; Fougereau, M.; Schiff, C.
Mol. Immunol. 31, 1099-1108, 1994
A:Title: The human pre-B cell receptor: structural constraints for a tentative model of
A:Reference number: I57832; MUID:95021318; PMID:7935499
A:Accession: I57832
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 1-145 <RES>
A:Cross-references: GB:S74019; NID:G693810; PIDN:AAB32118.1; PID:G693811
C:Genetics:
A:Gene: Vpre-B
A:Introns: 16/1
C:Superfamily: immunoglobulin V region; immunoglobulin homology

Query Match      38.6%; Score 207.5; DB 2; Length 145;
Best Local Similarity 48.8%; Pred. No. 2.8e-14;
Matches 42; Conservative 12; Mismatches 31; Indels 1; Gaps 1;

QY 9 GQVQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68
Db 34 GTTIRLCTLRNDH-DIGVSVYVYQQRPGHPFRFLRYFSQSDKSGQGPVPPRFSGSKD 92

QY 69 EAHNACVLTISPVQPEDDADYVCSVG 94
Db 93 VARNRGYLSISELQPEDEAVYCAVG 118

RESULT 6
L6HUST
IG lambda chain V-VI region (SUT) - human
C:Species: Homo sapiens (man)
C:Date: 30-Jun-1987 #sequence_revision 30-Jun-1987 #text_change 02-Sep-1997
C:Accession: A01988
R:Solomon, A.; Kyle, R.A.; Francione, B.
in Amyloidosis, Glenner, G.G., Osserman, E.F., Benditt, E.P., Calkins, E., Cohn, A.S., a
A:Title: Light chain variable region subgroups of monoclonal immunoglobulins in amyloid
A:Reference number: A01988
A:Accession: A01988
A:Molecule type: protein
A:Residues: 1-111 <SOL>
C:Genetics:
A:Gene: GDB:IGLV@
A:Cross-references: GDB:119342; OMIM:147240
A:Map position: 22q11.2-22q11.2
C:Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap
chain disulfide bonds. In some cases, such as IGA and IGM, the subunits associate into la
C:Superfamily: immunoglobulin V region; immunoglobulin homology
C:Keywords: heterotetramer; immunoglobulin
F:1-32/Region: framework 1
F:15-93/Domain: immunoglobulin homology <IMM>
F:23-35/Region: complementarity-determining 1
F:36-50/Region: framework 2
F:51-57/Region: complementarity-determining 2
F:58-91/Region: framework 3
F:92-100/Region: complementarity-determining 3
F:101-111/Region: framework 4
F:22-91/Disulfide bonds: #status predicted

Query Match      37.2%; Score 200; DB 1; Length 111;
Best Local Similarity 48.8%; Pred. No. 1.2e-13;
Matches 41; Conservative 12; Mismatches 25; Indels 6; Gaps 2;
```

QY 8 PGQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADIPDRSAAK 67  
Db 14 PGKTVITSCGSG--TIAGYVQWYQORPGRAPTVIPI-----EDTQPSGVDFRFGSI 67

QY 68 DEAHNACVLITSPVQPEDDADYYC 91  
Db 68 DRSSNSASLTISGLQTEDEADYYC 91

RESULT 7  
L6HUAR  
Ig lambda chain V-VI region (AR) - human (tentative sequence)  
C:Species: Homo sapiens (man)  
C:Date: 02-Apr-1982 #sequence\_revision 02-Apr-1982 #text\_change 31-Mar-2000  
C:Accession: A01987  
R:Stietten, K.; Natvig, J.B.; Husby, G.; Juul, J.  
Biochem. J. 195, 561-572, 1981  
A:Title: The complete amino acid sequence of a prototype immunoglobulin-lambda light-chain  
A:Reference number: A01987; MUID:82091000; PMID:6797401  
A:Contents: amyloid protein AR  
A:Accession: A01987  
A:Molecule type: protein  
A:Residues: 1-112 <SLE>  
A:Note: about half of the lambda chain C region is missing from this protein  
C:Comment: This protein was isolated from the spleen of a patient with amyloidosis.  
C:Genetics:  
A:Gene: GDB:IGLV@  
A:Cross-references: GDB:119342; OXIM:147240  
A:Map position: 22q11.2-22q11.2  
C:Complex: An immunoglobulin heterotetramer subunit consists of two identical light (lambda) chain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into larger complexes.  
C:Superfamily: immunoglobulin V region; immunoglobulin homology  
C:Keywords: amyloid; heterotetramer; immunoglobulin  
F:15-93/Domain: immunoglobulin homology <IMM>  
F:22-91/Disulfide bonds: #status predicted

Query Match 36.6%; Score 197; DB 1; Length 112;  
Best Local Similarity 47.6%; Pred. No. 2.6e-13;  
Matches 40; Conservative 15; Mismatches 23; Indels 6; Gaps 2;

QY 8 PGQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADIPDRSAAK 67  
Db 14 PGKTVITSCGSG--SIAGSFVQWYQORPGSAPTIVY-----DDNRPQGVDFRFGSI 67

QY 68 DEAHNACVLITSPVQPEDDADYYC 91  
Db 68 DDSANSASLTISGLKTEDEADYYC 91

RESULT 8  
S17399  
Ig lambda chain precursor - rabbit (fragment)  
C:Species: Oryctolagus cuniculus (domestic rabbit)  
C:Date: 19-Feb-1994 #sequence\_revision 10-Nov-1995 #text\_change 21-Jan-2000  
C:Accession: S17399  
R:Hayzer, D.J.; Young-Cooper, G.O.; Mage, R.G.; Jaton, J.C.  
Eur. J. Immunol. 20, 2707-2712, 1990  
A:Title: cDNA clones encoding immunoglobulin lambda chains from rabbit expressing the phage lambda display system  
A:Reference number: S17399; MUID:91099420; PMID:2125274  
A:Accession: S17399  
A:Status: preliminary  
A:Molecule type: mRNA  
A:Residues: 1-232 <HAY>  
A:Cross-references: EMBL:X57729; NID:gl593; PIDN:CAA40896.1; PID:gl594  
A:Note: the authors translated the codon TTA for residue 92 as Trp and AGC for residue 114 as Ser.  
C:Superfamily: immunoglobulin V region; immunoglobulin homology  
C:Keywords: heterotetramer; immunoglobulin  
F:147-215/Domain: immunoglobulin homology <IMM>

Query Match 35.8%; Score 192.5; DB 2; Length 232;  
Best Local Similarity 43.7%; Pred. No. 1.7e-12;  
Matches 38; Conservative 14; Mismatches 30; Indels 5; Gaps 2;

QY 9 GOVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68  
Db 30 GASAKLTCTLSAHT---YTIIDWYQQQGEAPRYLMEHKSDGSYTKGTGVPDRFSGSSS 86

QY 69 EAHNACVLITSPVQPEDDADYYCVGY 95  
Db 87 GADR--YLIIPSVQADDEADYYCGADY 111

RESULT 9  
S25755  
Ig lambda chain - human  
C:Species: Homo sapiens (man)  
C:Date: 22-Nov-1993 #sequence\_revision 26-May-1995 #text\_change 21-Jan-2000  
C:Accession: S25755  
R:Combratio, G.; Klobbeck, H.G.  
Eur. J. Immunol. 21, 1512-1522, 1991  
A:Title: V(lambda) and J(lambda)-C(lambda) gene segments of the human immunoglobulin lambda chain  
A:Reference number: S16439; MUID:91257162; PMID:1904362  
A:Accession: S25755  
A:Status: preliminary; translation not shown  
A:Molecule type: mRNA  
A:Residues: 1-243 <COM>  
A:Cross-references: EMBL:X57820; NID:g33739; PIDN:CAA40957.1; PID:g33740  
C:Superfamily: immunoglobulin V region; immunoglobulin homology  
C:Keywords: heterotetramer; immunoglobulin  
F:158-226/Domain: immunoglobulin homology <IMM>

Query Match 35.6%; Score 191.5; DB 2; Length 243;  
Best Local Similarity 41.1%; Pred. No. 2.2e-12;  
Matches 37; Conservative 19; Mismatches 33; Indels 1; Gaps 1;

QY 9 GOVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68  
Db 34 GASVLTCTLS--SGFSVGVDFWIRYQKPGNPRLYYHSDSNKSGGVESRFGSGND 92

QY 69 EAHNACVLITSPVQPEDDADYYCVGYGFS 98  
Db 93 ASANAGILIRISGLQLEVEADYYCGTWHNS 122

RESULT 10  
PS0055  
Ig lambda chain precursor V-II region - rabbit  
C:Species: Oryctolagus cuniculus (domestic rabbit)  
C:Date: 31-Mar-1990 #sequence\_revision 31-Mar-1990 #text\_change 23-Jul-1999  
C:Accession: PS0055  
R:Hayzer, D.J.; Jaton, J.C.  
Gene 80, 185-191, 1989  
A:Title: Cloning and sequencing of two functional rabbit germ-line immunoglobulin V lambda chain precursors  
A:Reference number: A91614; MUID:90006781; PMID:2507399  
A:Accession: PS0055  
A:Molecule type: DNA  
A:Residues: 1-120 <HAY>  
A:Cross-references: GB:M27840; NID:g341760; PIDN:AAA31363.1; PID:g552407  
A:Note: the authors translated the codon TTG for residue 97 as Trp  
C:Genetics: 17/1  
A:Introns: 17/1  
C:Superfamily: immunoglobulin V region; immunoglobulin homology  
C:Keywords: heterotetramer; immunoglobulin  
F:1-20/Domain: signal sequence #status predicted <SIG>  
F:21-120/Product: Ig lambda chain V-II region #status predicted <MAT>

Query Match 35.2%; Score 189.5; DB 2; Length 120;  
Best Local Similarity 43.7%; Pred. No. 1.6e-12;  
Matches 38; Conservative 13; Mismatches 31; Indels 5; Gaps 2;

QY 9 GOVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68  
Db 35 GASAKLTCTLSAHT---YTIIDWYQQQGEAPRYLMLKSDGSYTKGTGVPDRFSGSSS 91

QY 69 EAHNACVLITSPVQPEDDADYYCVGY 95





C;Superfamily: immunoglobulin V region; immunoglobulin homology  
C;Keywords: heterotetramer; immunoglobulin  
F;1-20/Domain: signal sequence #status predicted <SIG>  
F;21-120/Product: Ig lambda chain V-III region #status predicted <MAT>

Query Match 34.3%; Score 184.5; DB 2; Length 120;  
Best Local Similarity 44.7%; Pred. No. 5.3e-12;  
Matches 38; Conservative 13; Mismatches 29; Indels 5; Gaps 2;

QY 9 GQVAQLSCTLSPOHVTIRDVGSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAKD 68  
Db 35 GSAKLTCTLSAHKT--YYIEWYQQQGEAPRYLMQLKSDGSYTKGTGVPDRFSGSSS 91

QY 69 EAHNACVLITISPVQPEDDADYCSV 93  
Db 92 GADR--YLIISVQADDEADYICGV 114

RESULT 15  
S16848  
Ig lambda chain V-II region precursor - human  
C;Species: Homo sapiens (man)  
C;Date: 20-Feb-1995 #sequence\_revision 20-Feb-1995 #text\_change 21-Jan-2000  
C;Accession: S60297; S16848  
R;Kueppers, R.; Fischer, U.; Rajewsky, K.; Gause, A.  
Immunol. Lett. 34, 57-62, 1992  
A;Title: Immunoglobulin heavy and light chain gene sequences of a human CD5 positive imm  
A;Reference number: S60295; MUID:93122853; PMID:1282498  
A;Accession: S60297  
A;Status: Preliminary  
A;Molecule type: DNA  
A;Residues: 1-136 <KU2>  
A;Cross-references: EMBL:X62125; NID:938334; PIDN:CAA44056.1; PID:938335  
C;Superfamily: immunoglobulin V region; immunoglobulin homology  
C;Keywords: heterotetramer; immunoglobulin  
F;34-111/Domain: immunoglobulin homology <IMM>

Query Match 34.3%; Score 184.5; DB 2; Length 136;  
Best Local Similarity 45.9%; Pred. No. 6.1e-12;  
Matches 39; Conservative 14; Mismatches 25; Indels 7; Gaps 3;

QY 8 PGQVAQLSCTLSPOHVTIRDVGSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAK 67  
Db 33 PGQSVTISCTGTSSDVGAVNY-VSWYQHHPGKAPKLMYEVSE---RPSGVPDRFSGSK 87

QY 68 DEAHNACVLITISPVQPEDDADYCS 92  
Db 88 --SGNTASLTIVSLQAEDEADYICT 110

Search completed: June 28, 2004, 08:29:57  
Job time : 12.0405 secs

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GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:35 ; Search time 8.91892 Seconds  
(without alignments)  
577.979 Million cell updates/sec

Title: US-09-981-876-200\_COPY\_25\_123  
Perfect score: 538  
Sequence: 1 LDALLVFPQVAGLSTLSP.....PVQPEDDADYCVSGYGFSP 99

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 141681 seqs, 52070155 residues

Total number of hits satisfying chosen parameters: 141681

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : SwissProt\_42:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	538	100.0	123	1 VPR3_HUMAN	Q9UK13
2	233.5	43.4	142	1 VPR2_MOUSE	P13373 mus musculus
3	239.5	42.7	142	1 VPR1_MOUSE	P13372 mus musculus
4	207.5	38.6	145	1 VPRE_HUMAN	P12018 homo sapien
5	200	37.2	111	1 LV6C_HUMAN	P06317 homo sapien
6	197	36.6	112	1 LV6A_HUMAN	P01721 homo sapien
7	185.5	34.5	111	1 LV2F_HUMAN	P01709 homo sapien
8	185	34.4	111	1 LV6E_HUMAN	P06318 homo sapien
9	183	34.0	131	1 LV6D_HUMAN	P06319 homo sapien
10	182.5	33.9	111	1 LV2L_HUMAN	P80422 homo sapien
11	180.5	33.6	111	1 LV2G_HUMAN	P04209 homo sapien
12	175.5	32.6	112	1 LV2K_HUMAN	P01712 homo sapien
13	175	32.5	111	1 LV2I_HUMAN	P01710 homo sapien
14	172.5	32.1	109	1 LV2E_HUMAN	P01708 homo sapien
15	171	31.8	108	1 LV3A_HUMAN	P01714 homo sapien
16	167.5	31.1	110	1 LV2J_HUMAN	P01713 homo sapien
17	167.5	31.1	111	1 LV1D_HUMAN	P01702 homo sapien
18	167	31.0	130	1 LV1G_HUMAN	P06316 homo sapien
19	165	30.7	108	1 LV5A_HUMAN	P01719 homo sapien
20	165	30.7	109	1 LV1F_HUMAN	P04208 homo sapien
21	164	30.5	111	1 LV3B_HUMAN	P08048 homo sapien
22	163.5	30.4	111	1 LV2B_HUMAN	P01705 homo sapien
23	163.5	30.4	111	1 LV2H_HUMAN	P01711 homo sapien
24	162	30.1	106	1 LV4D_HUMAN	P01718 homo sapien
25	161	29.9	106	1 LV4B_HUMAN	P01716 homo sapien
26	161	29.9	109	1 KV3D_HUMAN	P01622 homo sapien
27	160	29.7	107	1 LV4C_HUMAN	P01717 homo sapien
28	160	29.7	108	1 KV3A_HUMAN	P01619 homo sapien
29	159.5	29.6	117	1 LV0A_HUMAN	P04211 homo sapien
30	158	29.4	106	1 LV4A_HUMAN	P01715 homo sapien
31	158	29.4	129	1 KV3L_HUMAN	P18135 homo sapien
32	156	29.0	109	1 KV3B_HUMAN	P01620 homo sapien
33	156	29.0	109	1 KV3G_HUMAN	P04206 homo sapien

34	155.5	28.9	112	1 LV1H_HUMAN	P06887 homo sapien
35	154.5	28.7	111	1 LV2A_HUMAN	P01704 homo sapien
36	154.5	28.7	111	1 LV2C_HUMAN	P01706 homo sapien
37	154.5	28.7	112	1 LV1B_HUMAN	P01700 homo sapien
38	153.5	28.5	111	1 LV2D_HUMAN	P01707 homo sapien
39	153	28.4	111	1 LV1C_HUMAN	P01701 homo sapien
40	152	28.3	112	1 LV6B_HUMAN	P01722 homo sapien
41	152	28.3	129	1 KV3M_HUMAN	P18136 homo sapien
42	148	27.5	100	1 KV3C_HUMAN	P01621 homo sapien
43	147.5	27.4	109	1 KV3E_HUMAN	P01623 homo sapien
44	147	27.3	109	1 LV1I_HUMAN	P06888 homo sapien
45	146	27.1	106	1 LV4E_HUMAN	P06889 homo sapien

ALIGNMENTS

RESULT 1  
VPR3\_HUMAN  
ID VPR3\_HUMAN STANDARD; PRT; 123 AA.  
AC Q9UK13;  
DT 16-OCT-2001 (Rel. 40, Created)  
DT 16-OCT-2001 (Rel. 40, Last sequence update)  
DT 15-MAR-2004 (Rel. 43, Last annotation update)  
DE Pre-B lymphocyte protein 3 precursor (VpreB3 protein) (N27C7-2).  
GN VPREB3.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=20169186; PubMed=10702669;  
RT Rosnet O., Mattei M.-G., Delattre O., Schiff C.;  
RA VPREB3: cDNA characterization and expression in human and chromosome  
mapping in human and mouse.  
RL Cytogenet. Cell Genet. 87:205-208(1999).  
RN [2]  
RP SEQUENCE FROM N.A.  
RA Shimizu N., Minosima S., Kawasaki K., Sasaki T., Hosono K.;  
RT "Molecular cloning of N27C7-2 gene."  
RL Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.  
RN [3]  
RP SEQUENCE FROM N.A.  
TX TISSUE=Testis;  
MEDLINE=22388257; PubMed=12477932;  
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Schuler G.D.,  
Klausner R.D., Collins P.S., Wagner L., Shenmen C.M., Schaefer C.F., Bhat N.K.,  
Altschul S.F., Zeeberg B., Buetow K.H., Wang J., Wang J., Hsieh F.,  
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,  
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,  
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,  
Brownstein M.J., Usdin T.B., Teshiyuki S., Carninci P., Prange C.,  
Rana S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,  
Bosak S.A., McSwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,  
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,  
Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,  
Rahy J., Helton E., Kettelman M., Madan A., Rodrigues S., Sanchez A.,  
Whiting M., Madan A., Young A.C., Shevchenko V., Bouffard G.G.,  
Brakesley A.C., Touchman J.W., Green E.D., Dickson M.C.,  
Rodriguez R.C., Grimwood J., Schmutz J., Myers R.M.,  
Rutterfield A.S.N., Krzywinski M.I., Skalska U., Smallos D.E.,  
Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;  
RT "Generation and initial analysis of more than 15,000 full-length  
human and mouse cDNA sequences";  
RL Proc Natl Acad Sci U S A. 99:16899-16903(2002).  
CC -!- FUNCTION: ASSOCIATES WITH THE IG-WU CHAIN TO FORM A MOLECULAR  
COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS.  
CC -!- TISSUE SPECIFICITY: Expressed in B cell precursors. Expressed in  
fetal liver, bone marrow, spleen and lymph node.  
CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.  
CC -!- SIMILARITY: Contains 1 immunoglobulin-like domain.

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 CC -----

DR EMBL; AF163825; AAF09451.1; -;  
 DR EMBL; AB050772; BAB83034.1; -;  
 DR EMBL; BC020666; AAH20666.1; -;  
 DR HSSP; P01709; 2MCG.  
 DR Genew; HGNC:12710; VPREB3..  
 DR MIM; 605017; -;  
 DR InterPro; IPR007110; IG-like.  
 DR InterPro; IPR003596; IG\_V.  
 DR Pfam; PF00047; IG; 1.  
 DR SMART; SM00406; IGV; 1.  
 DR PROSITE; PS0835; IG LIKE; 1.  
 KW Immunoglobulin domain; Signal.  
 FT SIGNAL 1 20 POTENTIAL.  
 FT CHAIN 21 123 PRE-B LYMPHOCYTE PROTEIN 3.  
 FT DOMAIN 21 123 IG-LIKE.  
 FT DISULFID 40 115 BY SIMILARITY.  
 SQ SEQUENCE 123 AA; 13710 MW; BF09AC5196059E85 CRC64;

Query Match 100.0%; Score 538; DB 1; Length 123;  
 Best Local Similarity 100.0%; Pred. No. 2.5e-52;  
 Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LDALLVFPQVAQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIP 60  
 DB 25 LDALLVFPQVAQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIP 84  
 QY 61 DRPSAAKDEAHNACVLITSPQEDDADYCVSGYGFSP 99  
 DB 85 DRPSAAKDEAHNACVLITSPQEDDADYCVSGYGFSP 123

RESULT 2  
 VPR2\_MOUSE  
 ID\_VPR2\_MOUSE STANDARD; PRT; 142 AA.  
 AC P13373;  
 DT 01-JAN-1990 (Rel. 13, Created)  
 DT 01-JAN-1990 (Rel. 13, Last sequence update)  
 DT 15-MAR-2004 (Rel. 43, Last annotation update)  
 DE Immunoglobulin omega chain precursor (VpreB2 protein).  
 GN VPREB2.  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=C57BL/6 X DBA/2J;  
 RX MEDLINE=88029315; PubMed=3117530;  
 RA Kudo A., Melchers F.;  
 RT "A second gene, VpreB in the lambda 5 locus of the mouse, which  
 RT appears to be selectively expressed in pre-B lymphocytes."  
 RL EMO J. 6:2267-2272(1987).  
 CC -!- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR  
 CC COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS. THIS  
 CC COMPLEX PRESUMABLY REGULATES IG GENE REARRANGEMENTS IN THE EARLY  
 CC STEPS OF B-CELL DIFFERENTIATION.  
 CC -!- TISSUE SPECIFICITY: ONLY EXPRESSED BY PRE-B-CELLS.  
 CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.

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 CC -----  
 DR EMBL; X05563; CAA29077.1; -;  
 DR PIR; B28344; B28344.  
 DR HSSP; P01607; 1RSI.  
 DR MGD; MGI:98937; Vpreb2.  
 DR InterPro; IPR007110; IG-like.  
 DR InterPro; IPR003596; IG\_V.  
 DR Pfam; PF00047; IG; 1.  
 DR SMART; SM00406; IGV; 1.  
 DR PROSITE; PS0835; IG LIKE; 1.  
 KW Immunoglobulin domain; Signal.  
 FT SIGNAL 1 19 POTENTIAL.  
 FT CHAIN 20 142 IMMUNOGLOBULIN OMEGA CHAIN.  
 FT DOMAIN 20 41 FRAMEWORK-1.  
 FT DOMAIN 42 56 COMPLEMENTARITY-DETERMINING-1.  
 FT DOMAIN 57 70 FRAMEWORK-2.  
 FT DOMAIN 71 81 COMPLEMENTARITY-DETERMINING-2.  
 FT DOMAIN 82 115 FRAMEWORK-3.  
 FT DISULFID 41 115 BY SIMILARITY.  
 SQ SEQUENCE 142 AA; 16052 MW; 7EA7128A4E63D920 CRC64;

Query Match 43.4%; Score 233.5; DB 1; Length 142;  
 Best Local Similarity 54.7%; Pred. No. 9.7e-19;  
 Matches 47; Conservative 9; Mismatches 29; Indels 1; Gaps 1;

QY 9 GQVAQLSCTLSFQHVITRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIPDRFSAKD 68  
 DB 34 GATIRLSCTLSNDH-NIGIYIYVYQQRPGHPRLRYFSHSDKHQGPDPFRFSGKD 92  
 QY 69 EARNACVLITSPQEDDADYCVSGV 94  
 DB 93 TARNLGYLSISELQPEDEAVYCAVG 118

RESULT 3  
 VPR1\_MOUSE  
 ID\_VPR1\_MOUSE STANDARD; PRT; 142 AA.  
 AC P13372;  
 DT 01-JAN-1990 (Rel. 13, Created)  
 DT 01-JAN-1990 (Rel. 13, Last sequence update)  
 DT 15-MAR-2004 (Rel. 43, Last annotation update)  
 DE Immunoglobulin Iota chain precursor (VpreB1 protein).  
 GN VPREB1.  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=C57BL/6 X DBA/2J;  
 RX MEDLINE=88029315; PubMed=3117530;  
 RA Kudo A., Melchers F.;  
 RT "A second gene, VpreB in the lambda 5 locus of the mouse, which  
 RT appears to be selectively expressed in pre-B lymphocytes."  
 RL EMO J. 6:2267-2272(1987).  
 CC -!- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR  
 CC COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS. THIS  
 CC COMPLEX PRESUMABLY REGULATES IG GENE REARRANGEMENTS IN THE EARLY  
 CC STEPS OF B-CELL DIFFERENTIATION.  
 CC -!- TISSUE SPECIFICITY: ONLY EXPRESSED BY PRE-B-CELLS.  
 CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.  
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 CC -----  
 DR EMBL; X05556; CAA29071.1; -;  
 DR EMBL; X05557; CAA29072.1; -;

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DR PIR; A28344.
DR HSSP; P01607; IREI.
DR MGD; MGI:98936; VpreB1.
DR GO; GO:0005886; C:plasma membrane; IPI.
DR GO; GO:0004872; F:receptor activity; IPI.
DR GO; GO:0030097; P:hemopoiesis; IMP.
DR GO; GO:0006955; P:immune response; IPI.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003596; IG_v.
DR Pfam; PF00047; IG; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS50835; IG LIKE; 1.
KW Immunoglobulin domain; Signal.
FT SIGNAL 1 19 POTENTIAL.
FT CHAIN 20 142 IMMUNOGLOBULIN IOTA CHAIN.
FT DOMAIN 20 41 FRAMEWORK-1.
FT DOMAIN 42 56 COMPLEMENTARITY-DETERMINING-1.
FT DOMAIN 57 70 FRAMEWORK-2.
FT DOMAIN 71 81 COMPLEMENTARITY-DETERMINING-2.
FT DOMAIN 82 115 FRAMEWORK-3.
FT DISULFID 41 115 BY SIMILARITY.
SQ SEQUENCE 142 AA; 16125 MW; 2E18BF963A0F448C CRC64;

Query Match 42.7%; Score 229.5; DB 1; Length 142;
Best Local Similarity 53.5%; Pred. No. 2.7e-18;
Matches 46; Conservative 9; Mismatches 30; Indels 1; Gaps 1;

QY 9 GOVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHRRPADIPDRFSAKD 68
Db 34 GATIRLSCTLSNDH-NIGYISYVYQORPGHPRFLRFSDHSHKXGQDIPRFRSGSKD 92

QY 69 EAHNACVLITSPVQPEDDADYCSVG 94
Db 93 TTRNLGYLSISELQPEDEAVYCAVG 118

RESULT 4
VPRE HUMAN STANDARD; PRT; 145 AA.
AC P12018;
DT 01-OCT-1999 (Rel. 12, Created)
DT 15-OCT-2001 (Rel. 40, Last sequence update)
DT 18-MAR-2004 (Rel. 43, Last annotation update)
DE Immunoglobulin iota chain precursor (V(pre)B protein) (VpreB protein)
DE (CD179a antigen).
GN VPREB1 OR VPREB.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Euthera; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
[1]
SEQUENCE FROM N.A.
MEDLINE=95021318; PubMed=7935499;
RA Guelpa-Fonlupt V., Bossy D., Alzari P., Fumoux F., Fougereau M.,
RA Schiff C.
RT "The human pre-B cell receptor: structural constraints for a tentative
RT model of the pseudo-light (psi L) chain."
RL Mol. Immunol. 31:1099-1108 (1994).
[2]
SEQUENCE FROM N.A.
MEDLINE=97228902; PubMed=9074928;
RA Kawasaki K., Minoshima S., Mine E., Shibuya K., Shintani A.,
RA Schmeits J.L., Wang J., Shimizu N.;
RT "One-megabase sequence analysis of the human immunoglobulin lambda
RT gene locus."
RL Genome Res. 7:250-261 (1997).
[3]
SEQUENCE OF 1-139 FROM N.A.
MEDLINE=88196069; PubMed=3258819;
RA Bauer S.R., Kudo A., Melchers F.;
RT "Structure and pre-B lymphocyte restricted expression of the VpreB in
RT humans and conservation of its structure in other mammalian
RT species.";
```

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RL PIR; 7:111-116 (1988).
CC -!- FUNCTION: ASSOCIATES WITH THE IG-MU CHAIN TO FORM A MOLECULAR
CC COMPLEX THAT IS EXPRESSED ON THE SURFACE OF PRE-B-CELLS. THIS
CC COMPLEX PRESUMABLY REGULATES IG GENE REARRANGEMENTS IN THE EARLY
CC STAGES OF B-CELL DIFFERENTIATION.
CC -!- SUBUNIT: Associates non-covalently with IGLL1.
CC -!- TISSUE SPECIFICITY: ONLY EXPRESSED BY PRE-B-CELLS.
CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.
CC -!- DATABASE: NAME=PROW; NOTE=PROW 1:59-63 (2000);
CC WWW="http://www.ncbi.nlm.nih.gov/prow/guide/574153212.g.htm".
CC
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CC
CC EMBL; D86992; BAA19987.1; -.
CC EMBL; D88270; BAA20030.1; -.
CC EMBL; S74019; AAB32118.1; -.
CC EMBL; M34927; AAB61292.1; -.
CC PIR; I57832; I57832.
CC HSSP; S00258; S00258.
CC PIR; S00258; S00258.
CC HSP; P80748; 2LOI.
CC GENE; HGNC:12709; VPREB1.
CC MIM; 605141; -.
CC GO; GO:0005576; C:extracellular; NAS.
CC GO; GO:0003823; F:antigen binding; NAS.
CC GO; GO:0006955; P:immune response; NAS.
CC InterPro; IPR007110; IG-like.
CC InterPro; IPR003596; IG_v.
CC Pfam; PF00047; IG; 1.
CC SMART; SM00406; IGV; 1.
CC PROSITE; PS50835; IG LIKE; 1.
KW Antigen; Signal; Immunoglobulin domain.
FT SIGNAL 1 19 POTENTIAL.
FT CHAIN 20 145 IMMUNOGLOBULIN IOTA CHAIN.
FT DOMAIN 20 41 FRAMEWORK-1.
FT DOMAIN 42 56 COMPLEMENTARITY-DETERMINING-1.
FT DOMAIN 57 70 FRAMEWORK-2.
FT DOMAIN 71 81 COMPLEMENTARITY-DETERMINING-2.
FT DOMAIN 82 115 FRAMEWORK-3.
FT DISULFID 41 115 BY SIMILARITY.
FT CONFLICT 10 10 L -> H (IN REF. 3). CRC64;
SQ SEQUENCE 145 AA; 16605 MW; 197665B13AF64D46 CRC64;

Query Match 38.6%; Score 207.5; DB 1; Length 145;
Best Local Similarity 48.8%; Pred. No. 7.2e-16;
Matches 42; Conservative 12; Mismatches 31; Indels 1; Gaps 1;

QY 9 GOVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHRRPADIPDRFSAKD 68
Db 34 GATIRLSCTLSNDH-NIGYISYVYQORPGHPRFLRFSDHSHKXGQDIPRFRSGSKD 92

QY 69 EAHNACVLITSPVQPEDDADYCSVG 94
Db 93 VARNRGYLSISELQPEDEAVYCAVG 118

RESULT 5
LV6C HUMAN STANDARD; PRT; 111 AA.
AC P06317;
DT 01-JAN-1988 (Rel. 06, Created)
DT 01-JAN-1988 (Rel. 06, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Ig lambda chain V-VI region SUT.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Euthera; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
```

RP SEQUENCE.  
RA Solomon A., Kyle R.A., Frangione B.;  
RT "Light chain variable region subgroups of monoclonal immunoglobulins  
in amyloidosis AL";  
RL (in) Glenner G.G., Osseman E.F., Benditt E.P., Calkins E.,  
Cohen A.S., Zucker-Franklin D. (eds.);  
RL Amyloidosis, pp.449-462, Plenum Press, New York (1986).  
DR PIR; A01988; L6HUST.  
DR PDB; 1CDG; 06-MAR-00.  
DR InterPro; IPR007110; Ig-like.  
DR InterPro; IPR003596; Ig\_V.  
DR Pfam; PF00047; Ig\_1.  
DR SMART; SM00406; IGV; 1.  
DR PROSITE; PS50835; IG LIKE; 1.  
KW Immunoglobulin V region, 3D-structure.  
FT DOMAIN 1 22  
FT DOMAIN 23 35  
FT DOMAIN 36 50  
FT DOMAIN 51 57  
FT DOMAIN 58 91  
FT DOMAIN 92 100  
FT DOMAIN 101 111  
FT DISULFID 122 91  
FT NON\_TER 111 111  
SQ SEQUENCE 111 AA; 12247 MW; 0941DD547D983598 CRC64;  
Query Match 37.2%; Score 200; DB 1; Length 111;  
Best Local Similarity 48.8%; Pred. No. 3.6e-15;  
Matches 41; Conservative 12; Mismatches 25; Indels 6; Gaps 2;  
QY 8 PGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAAK 67  
DB 14 PGKTVITFSGTGG--TIAGYVQWYQQRPGSAPTIVY---DDNQPSGVDPFRSGSI 67  
QY 68 DEAHNACVLITSPVQPEDDADYYC 91  
DB 68 DRSNSASLTISGLKTEDEADYYC 91  
RESULT 6  
LV6A\_HUMAN STANDARD; PRT; 112 AA.  
AC P01721;  
DT 21-JUL-1986 (Rel. 01, Created)  
DT 21-JUL-1986 (Rel. 01, Last sequence update)  
DT 10-OCT-2003 (Rel. 42, Last annotation update)  
DE Ig lambda chain V-VI region AR.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP X-RAY CRYSTALLOGRAPHY.  
RX MEDLINE=82091000; PubMed=6797401;  
RA Sletten K., Natvig J.B., Huseby G., Junl J.;  
RT "The complete amino acid sequence of a prototype  
immunoglobulin-lambda light-chain-type amyloid-fibril protein AR.";  
RL Biochem. J. 195;561-572(1981).  
CC -1- MISCELLANEOUS: ABOUT HALF OF THE LAMBDA CHAIN C REGION IS MISSING  
FROM THIS PROTEIN.  
CC -1- MISCELLANEOUS: THIS PROTEIN WAS ISOLATED FROM THE SPLEEN OF A  
PATIENT WITH AMYLOIDOSIS.  
CC -1- SIMILARITY: Contains 1 immunoglobulin-like domain.  
DR PIR; A01987; L6HUAR.  
DR HSP; P01709; ZMCG.  
DR GO; GO:0005576; C:extracellular; NAS.  
DR GO; GO:0003823; F:antigen binding; NAS.  
DR GO; GO:0006955; P:immune response; NAS.  
DR InterPro; IPR007110; Ig-like.  
DR Pfam; PF00047; Ig\_1.  
DR SMART; SM00406; IGV; 1.

DR PROSITE; PS50835; IG LIKE; 1.  
KW Immunoglobulin V region; Amyloid.  
FT DOMAIN 1 107  
FT NON\_TER 112 112  
SQ SEQUENCE 112 AA; 11918 MW; 570BCD9A368F1FE CRC64;  
Query Match 36.6%; Score 197; DB 1; Length 112;  
Best Local Similarity 47.6%; Pred. No. 7.7e-15;  
Matches 40; Conservative 15; Mismatches 23; Indels 6; Gaps 2;  
QY 8 PGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAAK 67  
DB 14 PGKTVITFSGTGG--SIADSFVQWYQQRPGSAPTIVY---DDNQPSGVDPFRSGSI 67  
QY 68 DEAHNACVLITSPVQPEDDADYYC 91  
DB 68 DRSNSASLTISGLKTEDEADYYC 91  
RESULT 7  
LV2P\_HUMAN STANDARD; PRT; 111 AA.  
AC P01709;  
DT 21-JUL-1986 (Rel. 01, Created)  
DT 21-JUL-1986 (Rel. 01, Last sequence update)  
DT 10-OCT-2003 (Rel. 42, Last annotation update)  
DE Ig lambda chain V-II region MGC.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=75013804; PubMed=4415202;  
RA Fett J.W., Deutsch H.F.;  
RT "Primary structure of the Mgc lambda chain.";  
RL Biochemistry 13:4102-4114(1974).  
RN [2]  
RP LAMBDA CHAIN GENES.  
RX MEDLINE=76093781; PubMed=812801;  
RA Fett J.W., Deutsch H.F.;  
RT "A new lambda-chain gene.";  
RL Immunochimistry 12:643-652(1975).  
RN [3]  
RP X-RAY CRYSTALLOGRAPHY (2.3 ANGSTROMS).  
RA Edmundson A.B., Ely K.R., Abola E.E., Schiffer M.,  
Panagiotopoulos N.;  
RT "Rotational allomerism and divergent evolution of domains in  
immunoglobulin light chains.";  
RL Biochemistry 14:3953-3961(1975).  
RN [4]  
RP X-RAY CRYSTALLOGRAPHY.  
RX MEDLINE=90133913; PubMed=2515285;  
RA Ely K.R., Herron J.N., Harker M., Edmundson A.B.;  
RT "Three-dimensional structure of a light chain dimer crystallized in  
water. Conformational flexibility of a molecule in two crystal  
forms.";  
RL J. Mol. Biol. 210:601-615(1989).  
CC -1- MISCELLANEOUS: This is a Bence-Jones protein.  
CC -1- MISCELLANEOUS: THE MCG-TYPE C REGION APPEARS TO BE CORRELATED WITH  
A VERY UNUSUAL V-REGION SUBSTITUTION, 103-THR ABOVE FOR GLY.  
CC SUGGESTING THAT THE V-C JOINING MECHANISM IS NOT ALWAYS RANDOM.  
CC -1- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE KERN+ AND MCG+  
MARKERS.  
CC -1- SIMILARITY: Contains 1 immunoglobulin-like domain.  
DR PIR; A90381; L2HUMC.  
DR PDB; 2MCG; 15-JUL-92.  
DR PDB; 1A8J; 17-JUN-98.  
DR PDB; 1DCL; 15-MAY-97.  
DR GO; GO:0005576; C:extracellular; NAS.  
DR GO; GO:0003823; F:antigen binding; NAS.  
DR GO; GO:0006955; P:immune response; NAS.  
DR InterPro; IPR007110; Ig-like.

[illegible]





```

RN      [1]
RP      SEQUENCE.
RX      MEDLINE=60006606; PubMed=113407;
RX      INFANTE A.J., Putnam F.W.;
RA      "Primary structure of a human IGA1 immunoglobulin. V. Amino acid
RT      sequence of a human IGA lambda light chain (Bur).";
RL      J. Biol. Chem. 254:9006-9016(1979).
RC      -!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE KERN* AND MCG*
CC      MARKERS.
CC      -!- MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM A MYELOMA PROTEIN.
CC      -!- SIMILARITY: Contains 1 immunoglobulin-like domain.
CC      PIR; A01974; L2HUBR.
DR      HSP; P01709; 2MCG.
DR      GO; GO:0005576; C:extracellular; NAS.
DR      GO; GO:0003823; F:antigen binding; NAS.
DR      GO; GO:0006955; P:immune response; NAS.
DR      InterPro; IPR007110; IG-like.
DR      InterPro; IPR003596; IG_V.
DR      Pfam; PF00047; IG: 1.
DR      SMART; SM00406; IGV; 1.
DR      PROSITE; PS50835; IG LIKE; 1.
KW      Immunoglobulin V region; Pyrrolidone carboxylic acid.
FT      DOMAIN 1 106 IG-LIKE.
FT      MOD RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
FT      DISULFID 22 90 BY SIMILARITY.
FT      SITE 91 91 APPEARS TO BE A FREE BUT UNREACTIVE
FT      SULPHYRYL GROUP.
FT      NON TER 109 109
SQ      SEQUENCE 109 AA; 11506 MW; BFD8AE1C5D267FAB CRC64;

Query Match 32.1%; Score 172.5; DB 1; Length 109;
Best Local Similarity 45.2%; Pred. No. 3,7e-12;
Matches 38; Conservative 12; Mismatches 27; Indels 7; Gaps 68

QY      8 PCQVQLSCTSPQGVTHRDYGVSWYQORAGSAPRVLVYRSEDHHRADIPDRESAAK 67
DB      14 PGHSVTISGIGTSNGVDYKY-VSWYQHPGKAPKLIY---EVSRSGVDPDRFSGK 68
QY      68 DEAHNAVLITSPVQPEDDADYVC 91
DB      69 --SGNTASLTISGLQAEDADYVC 90

RESULT 15
LV3A_HUMAN
ID      LV3A_HUMAN STANDARD; PRT; 108 AA.
AC      P01714;
DT      21-JUL-1986 (Rel. 01, Created)
DT      21-JUL-1986 (Rel. 01, Last sequence update)
DT      10-OCT-2003 (Rel. 42, Last annotation update)
DE      IG lambda chain V-III region SH.
OS      Homo sapiens (Human).
OC      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC      Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX      NCBI_TaxID=9606;
RN      [1]
RP      SEQUENCE.
RX      MEDLINE=70166723; PubMed=4909564;
RX      Titani K., Wikler M., Shinoda T., Putnam F.W.;
RA      "The amino acid sequence of a lambda type Bence-Jones protein. 3. The
RT      complete amino acid sequence and the location of the disulfide
RT      bridges.";
RL      J. Biol. Chem. 245:2171-2176(1970).
RC      -!- MISCELLANEOUS: This is a Bence-Jones protein.
CC      -!- SIMILARITY: Contains 1 immunoglobulin-like domain.
CC      PIR; A01980; L3HUSH.
DR      HSP; P80748; 2LQI.
DR      GO; GO:0005576; C:extracellular; NAS.
DR      GO; GO:0003823; F:antigen binding; NAS.
DR      GO; GO:0006955; P:immune response; NAS.
DR      InterPro; IPR007110; IG-like.
DR      InterPro; IPR003596; IG_V.
DR      Pfam; PF00047; IG: 1.

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```
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS50835; IG_LIKE; 1.
KW Immunoglobulin V region; Bence-Jones protein.
FT DOMAIN 1 97 IG-LIKE.
FT DISULFID 21 86
FT NON_TER 108 108
SQ SEQUENCE 108 AA; 11392 MW; E7E1229586411A56 CRC64;

Query Match      31.8%; Score 171; DB 1; Length 108;
Best Local Similarity 42.2%; Pred. No. 5.4e-12;
Matches 38; Conservative 17; Mismatches 25; Indels 10; Gaps 4;

Qy 3 ALLVFPQVQLSCTLSPOHVTIRDYGVSVMYQQFAGSAPRYLLYRSEEDHHPADIPDR 62
Db 8 AVSVALGTVRITC-----QGDSLRYDAWYQKRPQAPLLVIYGR---NNRPSGIPDR 59

Qy 63 FSAARDEAHNACVLITISVPQPEDDADYYCS 92
Db 60 FSGS-SSGHTAS-LTITGAQAEDEADYYCN 87
```

Search completed: June 28, 2004, 08:29:17  
Job time : 9.91892 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:34 ; Search time 32.5541 Seconds  
(without alignments)  
959.521 Million cell updates/sec

Title: US-09-981-876-200\_COPY\_25\_123  
Perfect score: 538  
Sequence: 1 LDALLVPPQVQLSCTLSP.....PVQPEDADYCSGVGVGFSFP 99

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1017041 seqs, 315518202 residues

Total number of hits satisfying chosen parameters: 1017041

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

SPTREMBL 25.\*

1: sp\_archaea.\*

2: sp\_bacteria.\*

3: sp\_fungi.\*

4: sp\_human.\*

5: sp\_invertebrate.\*

6: sp\_mammal.\*

7: sp\_mhc.\*

8: sp\_organelle.\*

9: sp\_phage.\*

10: sp\_plant.\*

11: sp\_rodent.\*

12: sp\_virus.\*

13: sp\_vertebrate.\*

14: sp\_unclassified.\*

15: sp\_rvirus.\*

16: sp\_bacteriap.\*

17: sp\_archaeap.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	372.5	69.2	123	11 Q61243	Q61243 mus musculus
2	215.5	40.1	230	4 Q722U3	Q722U3 homo sapien
3	199	37.0	135	4 Q9H5Z4	Q9H5Z4 homo sapien
4	197	36.6	112	4 Q96JD1	Q96JD1 homo sapien
5	193	35.9	112	4 Q96JD2	Q96JD2 homo sapien
6	189.5	35.2	116	4 Q96JD0	Q96JD0 homo sapien
7	175	32.5	237	4 Q9WTU6	Q9WTU6 homo sapien
8	174.5	32.4	236	4 Q96E61	Q96E61 homo sapien
9	172.5	32.1	237	4 Q9WUK4	Q9WUK4 homo sapien
10	172.5	32.1	240	4 Q9WUK3	Q9WUK3 homo sapien
11	169	31.4	107	4 Q9NSD6	Q9NSD6 homo sapien
12	166	30.9	109	4 Q9UL86	Q9UL86 homo sapien
13	165	30.7	233	4 Q9IBC9	Q9IBC9 homo sapien
14	161	29.9	234	4 Q9N355	Q9N355 homo sapien
15	161	29.9	235	11 Q99M11	Q99M11 mus musculus
16	160	29.7	108	4 Q96SBO	Q96SBO homo sapien

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17 158 29.4 109 4 Q9UL78
18 158 29.4 236 4 Q9NEU1
19 157 29.2 110 4 Q9TE83
20 156 29.0 81 4 Q7Z2E8
21 155 28.8 233 4 Q96I69
22 153 28.4 234 4 Q7Z2U7
23 151.5 28.2 100 6 Q776Z4
24 151 28.1 101 4 Q8I2D8
25 151 28.1 233 4 Q8N5F4
26 150.5 28.0 105 4 Q8WVJ6
27 148 27.5 107 4 Q9UL82
28 138 25.7 132 4 Q8TBD0
29 137 25.5 108 4 Q9UL83
30 134 24.9 97 4 Q43234
31 134 24.9 107 11 Q9ER29
32 133.5 24.6 484 11 Q8VEA0
33 132 24.5 109 4 Q9UL85
34 130.5 24.3 99 11 Q9JL74
35 129 24.0 111 11 Q8I1U6
36 128 23.8 107 4 Q96SA9
37 128 23.8 108 4 Q9UL77
38 127.5 23.7 239 4 Q8NEK0
39 127 23.6 108 4 Q9UL79
40 127 23.6 248 13 Q7SYU1
41 126 23.4 236 11 Q7TMK3
42 125 23.2 237 13 Q7SZ36
43 124.5 23.1 114 4 Q9UL80
44 124 23.0 131 11 Q811C3
45 124 23.0 494 4 Q96K68

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#### ALIGNMENTS

#### RESULT 1

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O61243 Q61243 PRELIMINARY; PRT; 123 AA.
ID AC Q61243;
DT 01-NOV-1996 (TRENBLrel. 01, Created)
DT 01-NOV-1996 (TRENBLrel. 01, Last sequence update)
DT 01-OCT-2003 (TRENBLrel. 25, Last annotation update)
DE SHS20 protein precursor (Pre-B lymphocyte gene 3).
GN VPREB3.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=BALB/C.
RX MEDLINE=93259124; PubMed=8491176;
RA Shirasawa T., Ohnishi K., Hagiwara S., Shigemoto K., Takabe Y.,
RA Rajewsky K., Takemori T.;
RT "A novel gene product associated with mu chains in immature B cells.";
RL EMBO J. 12:1827-1834(1993).
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/60; TISSUE=Stomach;
RX MEDLINE=21085660; PubMed=11217851;
RA Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gotohori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuehl P., Lewis S., Matsuo I., Nikaide I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombarts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,

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RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,  
 RA Suzuki H., Toyooka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,  
 RA Wyshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kontsuki S.,  
 RA Hayashizaki Y.,  
 RT "Functional annotation of a full-length mouse cDNA collection."  
 RL Nature 409:685-690(2001).  
 DR EMBL; D13208; BAA02495.1; -.  
 DR EMBL; AK008794; BAB25899.1; -.  
 DR PIR; S35302; S35302.  
 DR HSSP; P01709; 2MCG.  
 DR MGB; MGI:98936; vpreb3.  
 DR InterPro; IPR007110; IG-like.  
 DR InterPro; IPR003596; IG\_v.  
 DR Pfam; PF00047; IG\_1.  
 DR SMART; SM00406; IGV; 1.  
 DR PROSITE; PS50635; IG-LIKE; 1.  
 FT CHAIN 20 123  
 SQ SEQUENCE 123 AA; 13400 MW; 2AIAD371DICEE98F CRC64;  
 Query Match 69.2%; Score 372.5; DB 11; Length 123;  
 Best Local Similarity 69.4%; Pred. No. 6.5e-35;  
 Matches 68; Conservative 11; Mismatches 18; Indels 1; Gaps 1;  
 QY 2 DALLVFGVQVQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYVRSEEDHRRPADIPD 61  
 DB 27 DAFVFFGQDHLSCITNSQRTATGIGVSWYQQQGSAP-HLLYYAEHEHYRPADIPD 85  
 QY 62 RFSAKDEAHNACVLTISPVQEDDADYYCVGSGSP 99  
 DB 86 RFSAIVDAAHNACVLTISPVLPEDDADYFCIAHTPEP 123  
 RESULT 2  
 Q722U3  
 ID Q722U3 PRELIMINARY; PRT; 230 AA.  
 AC Q722U3  
 DT 01-OCT-2003 (TrEMBLrel. 25, Created)  
 DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)  
 DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)  
 DE Hypothetical protein.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 OX NCBITaxID=9606;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE=22388257; PubMed=12477932;  
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,  
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,  
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,  
 RA Hopkins R.F., Jordan H., Moore T., Max S.L., Wang J., Xsieh F.,  
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,  
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,  
 RA Brownstein M.J., Udén T.B., Toshiyuki S., Carninci P., Prange C.,  
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullany S.J.,  
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,  
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,  
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,  
 RA Fahey J., Helton E., Kettelman M., Madan A., Rodrigues S., Sanchez A.,  
 RA Whitting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,  
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,  
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,  
 RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,  
 RA Jones S.J., Marra M.A.;  
 RT "Generation and initial analysis of more than 15,000 full-length human  
 and mouse cDNA sequences."  
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).  
 RN [2]  
 RP SEQUENCE FROM N.A.  
 RA Strausberg R.;  
 RL Submitted (JUL-2003) to the EMBL/GenBank/DBJ databases.  
 DR EMBL; BC054893; AAH54893.1; -.  
 KW Hypothetical protein.

SQ SEQUENCE 230 AA; 24853 MW; 8BB60CC824BB886E CRC64;  
 Query Match 40.1%; Score 215.5; DB 4; Length 230;  
 Best Local Similarity 45.1%; Pred. No. 1.4e-16;  
 Matches 41; Conservative 20; Mismatches 29; Indels 1; Gaps 1;  
 QY 3 ALLVFPQVQVQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYVRSEEDHRRPADIPDR 62  
 DB 18 SLASPGASASLTCLR-RGFVYVDYRIYWTYQQSGRSPQYLLHRSDSDXQQSGVPSR 76  
 QY 63 FSAKDEAHNACVLTISPVQEDDADYYCVS 93  
 DB 77 FSGSKDASANAGIIVISGLRSEADYYCMV 107  
 RESULT 3  
 Q9H5Z4  
 ID Q9H5Z4 PRELIMINARY; PRT; 135 AA.  
 AC Q9H5Z4;  
 DT 01-MAR-2001 (TrEMBLrel. 16, Created)  
 DT 01-MAR-2001 (TrEMBLrel. 16, Last sequence update)  
 DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)  
 DE Hypothetical protein FLJ22755.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 OX NCBITaxID=9606;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=ileal mucosa;  
 RA Kawakami T., Noguchi S., Itoh T., Shigeta K., Senba T., Matsumura K.,  
 RA Nakajima Y., Mizuno T., Morinaga M., Tanigami A., Fujiwara T., Ono T.,  
 RA Yamada K., Fujii Y., Ozaki K., Hirao M., Ohmori Y., Ota T., Suzuki Y.,  
 RA Obayashi M., Nishi T., Shibahara T., Tanaka T., Nakamura Y.,  
 RA Isogai T., Sugano S.;  
 RT "NEDO human cDNA sequencing project";  
 RL Submitted (AUG-2000) to the EMBL/GenBank/DBJ databases.  
 DR EMBL; AK026408; BAB15473.1; -.  
 DR HSSP; P01607; IREI.  
 DR InterPro; IPR007110; IG-like.  
 DR InterPro; IPR003596; IG\_v.  
 DR SMART; SM00406; IGV; 1.  
 KW Hypothetical protein.  
 SQ SEQUENCE 135 AA; 14780 MW; 652492DED930F401 CRC64;  
 Query Match 37.0%; Score 199; DB 4; Length 135;  
 Best Local Similarity 45.3%; Pred. No. 5.9e-15;  
 Matches 34; Conservative 17; Mismatches 24; Indels 0; Gaps 0;  
 QY 24 TIRDYGVSWYQORAGSAPRYLLYVRSEEDHRRPADIPDRSAKDEAHNACVLTISPVQ 83  
 DB 7 SVGFQWIRWYQQKPGNPPRYLLYHSDNSKGGQSGVPSRFGSNDASANAGIIRISGLQP 66  
 QY 84 EDDADYYCVSGYGF 98  
 DB 67 EDEADYYCGTWHNS 81  
 RESULT 4  
 Q96JDI  
 ID Q96JDI PRELIMINARY; PRT; 112 AA.  
 AC Q96JDI;  
 DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
 DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
 DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)  
 DE Amyloid lambda 6 light chain variable region PIP (Fragment).  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 OX NCBITaxID=9606;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Bone marrow;

```
RA Perfetti V., Casarini S., Colli Vignarelli M., Merlini G.;
RT "Amyloid lambda 6 light chain variable region PIP.";
RL Submitted (MAY-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL: AF267874; AAK58586.1; -.
DR PIR: A30323; A30323.
DR InterPro: IPR007110; Ig-like.
DR InterPro: IPR003596; Ig_v.
DR Pfam: PF00047; Ig; 1.
DR SMART: SM00406; IGV; 1.
DR PROSITE: PS50835; IG_LIKE; 1.
FT NON_TER 1
FT NON_TER 112
SQ SEQUENCE 112 AA; 12047 MW; 0D3885AC23567B9F CRC64;

Query Match 36.6%; Score 197; DB 4; Length 112;
Best Local Similarity 47.6%; Pred. No. 8e-15;
Matches 40; Conservative 14; Mismatches 24; Indels 6; Gaps 2;

Qy 8 PGQVAQLSCTLSPOHVTIRDVGVSWYQORAGSAPRYLLYRSEEDHRRPADIPDRFSAK 67
Db 14 PGKTTISCTSSG--IASNSYQVQYQORPGSAPTIVY----EDNQRPSGVDRFSGSI 67
Qy 68 DEAHNACVLITSPVQPEDDADYYC 91
Db 68 DSSNSASLTISGLKTEDEADYYC 91

RESULT 5
Q96JD2 PRELIMINARY; PRT; 112 AA.
AC Q96JD2;
ID Q96JD2;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Amyloid lambda 6 light chain variable region NEG (fragment).
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP TISSUE=Bone marrow;
RC Perfetti V., Casarini S., Colli Vignarelli M., Merlini G.;
RT "Amyloid lambda 6 light chain variable region SAR.";
RL Submitted (MAY-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL: AF267875; AAK58587.1; -.
DR InterPro: IPR007110; Ig-like.
DR InterPro: IPR003596; Ig_v.
DR Pfam: PF00047; Ig; 1.
DR SMART: SM00406; IGV; 1.
DR PROSITE: PS50835; IG_LIKE; 1.
FT NON_TER 1
FT NON_TER 116
SQ SEQUENCE 116 AA; 12294 MW; F7B0E9F49FAE369E CRC64;

Query Match 35.2%; Score 189.5; DB 4; Length 116;
Best Local Similarity 44.6%; Pred. No. 6e-14;
Matches 41; Conservative 16; Mismatches 24; Indels 11; Gaps 4;

Qy 8 PGQVAQLSCTLSPOHVTIRDVGVSWYQORAGSAPRYLLYRSEEDHRRPADIPDRFSAK 67
Db 14 PGKTTISCTSSGSI-TNY-VQWYQLRFGSAPTIVY----EDNQRPSGVDRFSGSI 67
Qy 68 DEAHNACVLITSPVQPEDDADYYC-----SVG 94
Db 68 DSSNSASLTISGLKTEDEADYYCQSYDSSIG 99

RESULT 7
Q8WTU6 PRELIMINARY; PRT; 237 AA.
AC Q8WTU6;
ID Q8WTU6;
DT 01-MAR-2002 (TrEMBLrel. 20, Created)
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC Strausberg R.;
RT TISSUE=Tonsil;
RL Submitted (JAN-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL: BC022098; AAH22098.1; -.
DR PIR: S12441; S12441.
DR InterPro: IPR007110; Ig-like.
DR InterPro: IPR003006; Ig_MHC.
DR InterPro: IPR003596; Ig_v.
DR Pfam: PF00047; Ig; 2.
DR SMART: SM00406; IGV; 1.
DR PROSITE: PS50835; IG_LIKE; 2.
DR PROSITE: PS00290; IG_MHC; 1.
KW Hypothetical protein.
SQ SEQUENCE 237 AA; 24884 MW; B6CF371E753968E8 CRC64;

Query Match 32.5%; Score 175; DB 4; Length 237;
Best Local Similarity 42.9%; Pred. No. 6.5e-12;
Matches 39; Conservative 16; Mismatches 28; Indels 8; Gaps 4;

Qy 8 PGQVAQLSCTLSPOHVTIRDVGVSWYQORAGSAPRYLLYRSEEDHRRPADIPDRFSAK 67
Db 33 PGQVITISCTSSNIG-AGYDVHWYQLPGTAPKLLIYGN----NRPSGVDRFSGSK 87
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RP SEQUENCE FROM N.A.
RC TISSUE=Lymphocytes;
RA Hohmann A.;
RT "Autoimmunity.";
RL Submitted (JUL-1995) to the EMBL/GenBank/DBJ databases.
DR EMBL; L43092; AAA69746.2; -.
DR HSP; P01709; 2MCG.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS00835; IG_LIKE; 1.
FT NON_TER 1
FT NON_TER 107
SQ SEQUENCE 107 AA; 11306 MW; A2B04B37187A5F00 CRC64;

Query Match 31.4%; Score 169; DB 4; Length 107;
Best Local Similarity 41.7%; Pred. No. 1.2e-11;
Matches 35; Conservative 17; Mismatches 22; Indels 10; Gaps 3;

Qy 9 GQVAQLSCTLSPPQHVITRDYGVSWYQQRAGSAPRYLLYRSSEDHRRPADIPDRFSAAKD 68
Db 13 GQTVRIIC---QGSLSRYASWYQKPGQAPVLVIYK---NNRPSGIPDRFSGS-- 62

Qy 69 EAHNACVLITSPVQPEDDADYYCS 92
Db 63 SSGNTASLTITGAQAEDADYYCN 86

RESULT 12
Q9UL86 PRELIMINARY; PRT; 109 AA.
AC Q9UL86;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Myosin-reactive immunoglobulin kappa chain variable region
DE (Fragment);
OS Homo sapiens (Human);
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=98277139; PubMed=9614934;
RA Wu X., Liu B., Van der Werwe P.L., Kallis N.N., Berney S.M.,
RA Young D.C.;
RT "Myosin-reactive autoantibodies in rheumatic carditis and normal
RT fetus.";
RL Clin. Immunol. Immunopathol. 87:184-192 (1998).
DR EMBL; AF035028; AAD56264.1; -.
DR PIR; B30607; B30607.
DR PIR; I30601; I30601.
DR HSP; P80362; 1MTL.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS00835; IG_LIKE; 1.
FT NON_TER 1
FT NON_TER 109
SQ SEQUENCE 109 AA; 11928 MW; 243325F72C7DAC83 CRC64;

Query Match 30.9%; Score 166; DB 4; Length 109;
Best Local Similarity 44.2%; Pred. No. 2.8e-11;
Matches 42; Conservative 11; Mismatches 32; Indels 10; Gaps 4;

Qy 4 LLVFPQVAQLSCTLSPPQHVITRDYGVSWYQQRAGSAPRYLLYRSSEDHRRPADIPDRF 63
Db 11 LSLFPGERATLSQASQ---SVSSSYLAWYQKPGQAPRLIYGTSS----RATGIPDRF 63

Qy 64 SAAKDEAHNACVLITSPVQPEDDADYYCSVGYGS 98
Db 63 SSGNTASLTITGAQAEDADYYCN 86
```

```
Db 64 SGSGSETD--FTLTISRLPEDEFAVYQC-QYGSS 95

RESULT 13
Q8TBC9 PRELIMINARY; PRT; 233 AA.
AC Q8TBC9;
DT 01-JUN-2002 (TrEMBLrel. 21, Created)
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human);
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=B-cell;
RA Strausberg R.;
RL Submitted (FEB-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC022823; AAH22823.1; -.
DR PIR; S12442; S12442.
DR PIR; S30526; S30526.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003006; Ig_MHC.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 2.
DR SMART; SM00406; IGV; 1.
DR PROSITE; PS00835; IG_LIKE; 2.
DR PROSITE; PS00290; IG_MHC; 1.
KW Hypothetical protein.
SQ SEQUENCE 233 AA; 24867 MW; 367411BFD6F4DF92 CRC64;

Query Match 30.7%; Score 165; DB 4; Length 233;
Best Local Similarity 41.6%; Pred. No. 8.9e-11;
Matches 37; Conservative 15; Mismatches 21; Indels 16; Gaps 4;

Qy 6 VFPQVAQLSCT---LSPQHVITRDYGVSWYQQRAGSAPRYLLYRSSEDHRRPADIPDR 62
Db 31 VFPQTVARITCSGDALPKQY-----AYWQKPGQAPVLVIY---KDNERPSPGIPER 79

Qy 63 FSAKDEAHNACVLITSPVQPEDDADYYC 91
Db 80 FSGS--SSGTTVTLTISGVQAEADADYYC 106

RESULT 14
Q8N355 PRELIMINARY; PRT; 234 AA.
AC Q8N355;
DT 01-OCT-2002 (TrEMBLrel. 22, Created)
DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human);
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Brain;
RA Strausberg R.;
RL Submitted (APR-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC028090; AAH28090.1; -.
DR PIR; S12441; S12441.
DR InterPro; IPR003599; Ig.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003597; Ig_ci.
DR InterPro; IPR003006; Ig_MHC.
DR InterPro; IPR003596; Ig_v.
DR Pfam; PF00047; Ig; 2.
DR SMART; SM00409; IG; 2.
DR SMART; SM00407; IGL1; 1.
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GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: June 28, 2004, 08:24:34 ; Search time 45.4865 Seconds  
(without alignments)  
614.956 Million cell updates/sec

Title: US-09-981-876-200\_COPY\_25\_123

Perfect score: 538

Sequence: 1 LDALLVPGQVAQLSCTLSP.....PVPQEDADYCVSGYGFSP 99

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1586107 seqs, 282547505 residues

Total number of hits satisfying chosen parameters: 1586107

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : A\_Geneseq\_29Jan04:\*

- 1: Geneseqp1980s:\*
- 2: Geneseqp1990s:\*
- 3: Geneseqp2000s:\*
- 4: Geneseqp2001s:\*
- 5: Geneseqp2002s:\*
- 6: Geneseqp2003as:\*
- 7: Geneseqp2003bs:\*
- 8: Geneseqp2004s:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	538	100.0	113	4	AAM41476 Human pol
2	538	100.0	123	2	AAM41476 Human pol
3	538	100.0	123	3	AAY6655 Membrane-
4	538	100.0	123	3	AAB24061 Human PRO
5	538	100.0	123	4	AAU12372 Human PRO
6	538	100.0	123	4	AAU12372 Human PRO
7	538	100.0	123	6	ABU57993 Human PRO
8	538	100.0	123	6	ABU57993 Human PRO
9	538	100.0	123	6	ABU57993 Human PRO
10	538	100.0	123	6	ABU57993 Human PRO
11	538	100.0	123	6	ABU57993 Human PRO
12	538	100.0	123	6	ABU57993 Human PRO
13	538	100.0	123	6	ABU57993 Human PRO
14	538	100.0	123	6	ABU57993 Human PRO
15	538	100.0	123	6	ABU57993 Human PRO
16	538	100.0	123	6	ABU57993 Human PRO
17	538	100.0	123	6	ABU57993 Human PRO
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20	538	100.0	123	6	ABU57993 Human PRO
21	538	100.0	123	6	ABU57993 Human PRO
22	538	100.0	123	6	ABU57993 Human PRO
23	538	100.0	123	6	ABU57993 Human PRO
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27	538	100.0	123	6	ABU10839 Human PRO
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29	538	100.0	123	6	ABU10839 Human PRO
30	538	100.0	123	6	ABU10839 Human PRO
31	538	100.0	123	6	ABU10839 Human PRO
32	538	100.0	123	6	ABU10839 Human PRO
33	538	100.0	123	6	ABU10839 Human PRO
34	538	100.0	123	6	ABU10839 Human PRO
35	538	100.0	123	6	ABU10839 Human PRO
36	538	100.0	123	6	ABU10839 Human PRO
37	538	100.0	123	6	ABU10839 Human PRO
38	538	100.0	123	6	ABU10839 Human PRO
39	538	100.0	123	6	ABU10839 Human PRO
40	538	100.0	123	6	ABU10839 Human PRO
41	538	100.0	123	6	ABU10839 Human PRO
42	538	100.0	123	6	ABU10839 Human PRO
43	538	100.0	123	6	ABU10839 Human PRO
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#### ALIGNMENTS

RESULT 1  
AAM41476  
ID AAM41476 standard; protein; 113 AA.  
XX  
AC AAM41476;  
XX  
DT 22-OCT-2001 (first entry)  
XX  
DE Human polypeptide SEQ ID NO 6407.  
XX  
KW Human; neotropic; immunosuppressant; cytostatic; gene therapy; cancer;  
KW peripheral nervous system; neuropathy; central nervous system; CNS;  
KW Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic;  
KW amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;  
KW chemokine; thrombolytic; drug screening; arthritis; inflammation;  
KW leukaemia.  
XX  
CS Homo sapiens.  
XX  
PN WO200153312-A1.  
XX  
PD 26-JUL-2001.  
XX  
PF 26-DEC-2000; 2000WO-US034263.  
XX  
PR 23-DEC-1999; 99US-00471275.  
XX  
PR 21-JAN-2000; 2000US-00488725.  
XX  
PR 25-APR-2000; 2000US-00552317.  
XX  
PR 20-JUN-2000; 2000US-00598042.  
XX  
PR 19-JUL-2000; 2000US-00620312.  
XX  
PR 03-AUG-2000; 2000US-00653450.  
XX  
PR 14-SEP-2000; 2000US-00662191.  
XX  
PR 19-OCT-2000; 2000US-00693036.  
XX  
PR 29-NOV-2000; 2000US-00727344.  
XX  
(HYSE-) HYSEQ INC.  
XX  
PI Tang YT, Liu C, Asundi V, Chen R, Ma Y, Qian XB, Ren F, Wang D;  
PI Wang J, Wang Z, Wehrman T, Xu C, Xue AJ, Yang Y, Zhang J, Zhao QA;  
PI Zhou P, Goodrich R, Drmanac RT;  
XX  
DR WPI; 2001-442253/47.  
XX  
DR N-PSDB; AAI60632.  
XX  
XX Novel nucleic acids and polypeptides, useful for treating disorders such  
XX as central nervous system injuries.  
XX  
PS Example 2; SEQ ID NO 6407; 10078pp; English.

XX The invention relates to human nucleic acids (AA157798-AA161369) and the  
CC encoded polypeptides (AA135642-AA42213) with nootropic  
CC immunosuppressant and cytostatic activity. The polynucleotides are useful  
CC in gene therapy. A composition containing a polypeptide or polynucleotide  
CC of the invention may be used to treat diseases of the peripheral nervous  
CC system, such as peripheral nervous injuries, peripheral neuropathy and  
CC localised neuropathies and central nervous system diseases, such as  
CC Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic  
CC lateral sclerosis, and Shy-Drager Syndrome. Other uses include the  
CC utilisation of the activities such as: Immune system suppression,  
CC Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic  
CC and thrombolytic activity, cancer diagnosis and therapy, drug screening,  
CC assays for receptor activity, arthritis and inflammation, leukaemias and  
CC C.N.S disorders. Note: The sequence data for this patent did not form  
CC part of the printed specification  
XX  
XX Sequence 113 AA;  
  
Query Match 100.0%; Score 538; DB 4; Length 113;  
Best Local Similarity 100.0%; Pred. No. 9.1e-52;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
Qy 1 LDALLVFGQVAQSCSLSPQHVIRDYGVSWYQQRAGSAPRYLLYYRSEEDHRRADIP 60  
Db 15 LDALLVFGQVAQSCSLSPQHVIRDYGVSWYQQRAGSAPRYLLYYRSEEDHRRADIP 74  
  
Qy 61 DRFSAKDEAHNACVLTISPVQPEDDADYXCVGYGFSP 99  
Db 75 DRFSAKDEAHNACVLTISPVQPEDDADYXCVGYGFSP 113  
  
RESULT 2  
AAW75123  
ID AAW75123 standard; protein; 123 AA.  
XX AC AAW75123;  
XX  
DT 25-MAR-2003 (revised)  
DT 28-JAN-1999 (first entry)  
XX  
DE Human secreted protein encoded by gene 67 clone HRGDF73.  
XX  
KW Human; secreted protein; fusion protein; gene therapy; protein therapy;  
KW diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia;  
KW developmental abnormality; foetal deficiency; blood; allergy; renal;  
KW immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma;  
KW inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;  
KW cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus;  
KW osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;  
KW endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.  
XX  
OS Homo sapiens.  
XX  
XX WO9839446-A2.  
XX  
PD 11-SEP-1998.  
XX  
PF 06-MAR-1998; 98WO-US004482.  
XX  
PR 07-MAR-1997; 97US-0038621P.  
PR 07-MAR-1997; 97US-0040161P.  
PR 07-MAR-1997; 97US-0040162P.  
PR 07-MAR-1997; 97US-0040163P.  
PR 07-MAR-1997; 97US-0040333P.  
PR 07-MAR-1997; 97US-0040334P.  
PR 07-MAR-1997; 97US-0040336P.  
PR 07-MAR-1997; 97US-0040626P.  
PR 11-APR-1997; 97US-0043311P.  
PR 11-APR-1997; 97US-0043312P.  
PR 11-APR-1997; 97US-0043313P.  
PR 11-APR-1997; 97US-0043314P.  
PR 11-APR-1997; 97US-0043315P.  
  
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PR 11-APR-1997; 97US-0043569P.  
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PR 11-APR-1997; 97US-0043672P.  
PR 11-APR-1997; 97US-0043674P.  
PR 23-MAY-1997; 97US-0047492P.  
PR 23-MAY-1997; 97US-0047500P.  
PR 23-MAY-1997; 97US-0047501P.  
PR 23-MAY-1997; 97US-0047502P.  
PR 23-MAY-1997; 97US-0047503P.  
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PR 23-MAY-1997; 97US-0047582P.  
PR 23-MAY-1997; 97US-0047583P.  
PR 23-MAY-1997; 97US-0047584P.  
PR 23-MAY-1997; 97US-0047585P.  
PR 23-MAY-1997; 97US-0047586P.  
PR 23-MAY-1997; 97US-0047587P.  
PR 23-MAY-1997; 97US-0047588P.  
PR 23-MAY-1997; 97US-0047589P.  
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PR 23-MAY-1997; 97US-0047592P.  
PR 23-MAY-1997; 97US-0047593P.  
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PR 23-MAY-1997; 97US-0047615P.  
PR 23-MAY-1997; 97US-0047617P.  
PR 23-MAY-1997; 97US-0047618P.  
PR 23-MAY-1997; 97US-0047632P.  
PR 23-MAY-1997; 97US-0047633P.  
PR 06-JUN-1997; 97US-0048964P.  
PR 06-JUN-1997; 97US-0048974P.  
PR 22-AUG-1997; 97US-0056630P.  
PR 22-AUG-1997; 97US-0056631P.  
PR 22-AUG-1997; 97US-0056632P.  
PR 22-AUG-1997; 97US-0056636P.  
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PR 22-AUG-1997; 97US-0056875P.  
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PR 22-AUG-1997; 97US-0056877P.  
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PR 22-AUG-1997; 97US-0056881P.  
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PR 22-AUG-1997; 97US-0056886P.  
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PR 22-AUG-1997; 97US-0056893P.  
PR 22-AUG-1997; 97US-0056894P.

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PR 22-AUG-1997; 97US-0056903P.
PR 22-AUG-1997; 97US-0056908P.
PR 22-AUG-1997; 97US-0056909P.
PR 22-AUG-1997; 97US-0056910P.
PR 22-AUG-1997; 97US-0056911P.
PR 05-SEP-1997; 97US-0057650P.
PR 03-SEP-1997; 97US-0057761P.
XX
XX (HUMA-) HUMAN GENOME SCI INC.
XX Ruben SM, Rosen CA, Fischer CL, Soppet DR, Carter KC;
PI Bednarek DP, Endress GA, Yu G, Ni J, Feng P, Young PE, Greene JM;
PI Farrie AM, Duan R, Hu J, Florence KA, Olsen HS, Ebner R, Brewer LA;
PI Moore PA, Shi Y, Lafleur DW, Li Y, Zeng Z, Kyaw H;
XX WPI; 1998-609887/51.
DR N-PSDE; AAV34220.
XX
XX New isolated human genes and the secreted polypeptides they encode -
PT useful for diagnosis and treatment of e.g. cancers, neurological
PT disorders, immune diseases, inflammation or blood disorders.
XX
PS Claim 1; Page 320-321; 447pp; English.
XX
XX This sequence represents a secreted human protein encoded by the gene
CC clone detailed in the descriptor line. The gene can be used to generate
CC fusion proteins by linking to the gene to a human immunoglobulin Fc
CC portion (e.g. AAV34145) for increasing the stability of the fused protein
CC as compared to the human protein only. The invention relates to 70 novel
CC genes and their fragments (nucleic acid sequences: AAV34154-V34276; amino
CC acid sequences AAV75057-W75179) which are useful for preventing, treating
CC or ameliorating medical conditions e.g. by protein or gene therapy. Also,
CC pathological conditions can be diagnosed by determining the amount of the
CC new polypeptides in a sample or by determining the presence of mutations
CC in the new polynucleotides. Specific uses are described for each of the
CC 70 polynucleotides, based on which tissues they are most highly expressed
CC in (see AAV34154 for described uses). (Updated on 25-MAR-2003 to correct
CC PF field.) (Updated on 25-MAR-2003 to correct PI field.)
XX
XX Sequence 123 AA;
SQ
Query Match 100.0%; Score 538; DB 2; Length 123;
Best Local Similarity 100.0%; Pred. No. 1e-51;
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 LDALLVPGQVAQLSCTLSPOHVTIRDYGVSQYQVQAGSAPRYLLYYRSEDDHRRPADIP 60
Db 25 LDALLVPGQVAQLSCTLSPOHVTIRDYGVSQYQVQAGSAPRYLLYYRSEDDHRRPADIP 84
QY 61 DRFSAKDEAHNACVLTISPQVEDDADYCVSGYGFSP 99
Db 85 DRFSAKDEAHNACVLTISPQVEDDADYCVSGYGFSP 123
RESULT 3
AAV66655
ID AAV66655 standard; protein; 123 AA.
XX
XX AAV66655;
XX
XX 05-APR-2000 (first entry)
XX
XX Membrane-bound protein PRO619.
DE
XX Membrane-bound polypeptide; PRO polypeptide; LDL receptor; TIE ligand;
KW pharmaceutical; receptor immunoadhesin; gene mapping.
XX Homo sapiens.
XX
XX WO9963088-A2.
XX
XX 09-DEC-1999.
XX
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PP 02-JUN-1999; 99WO-US012252.
XX
XX 02-JUN-1998; 98US-0087607P.
PR 02-JUN-1998; 98US-0087609P.
PR 02-JUN-1998; 98US-0087759P.
PR 03-JUN-1998; 98US-0087827P.
PR 04-JUN-1998; 98US-0088021P.
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PR 09-JUN-1998; 98US-0088655P.
PR 10-JUN-1998; 98US-0088722P.
PR 10-JUN-1998; 98US-0088730P.
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PR 10-JUN-1998; 98US-0088824P.
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PR 11-JUN-1998; 98US-0088876P.
PR 12-JUN-1998; 98US-0089090P.
PR 12-JUN-1998; 98US-0089105P.
PR 16-JUN-1998; 98US-0089340P.
PR 16-JUN-1998; 98US-0089512P.
PR 16-JUN-1998; 98US-0089514P.
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PR 17-JUN-1998; 98US-0089653P.
PR 18-JUN-1998; 98US-0089801P.
PR 18-JUN-1998; 98US-0089907P.
PR 18-JUN-1998; 98US-0089908P.
PR 19-JUN-1998; 98US-0089947P.
PR 19-JUN-1998; 98US-0089948P.
PR 19-JUN-1998; 98US-0089952P.
PR 22-JUN-1998; 98US-0090246P.
PR 22-JUN-1998; 98US-0090252P.
PR 22-JUN-1998; 98US-0090254P.
PR 23-JUN-1998; 98US-0090349P.
PR 23-JUN-1998; 98US-0090355P.
PR 24-JUN-1998; 98US-0090429P.
PR 24-JUN-1998; 98US-0090431P.
PR 24-JUN-1998; 98US-0090435P.
PR 24-JUN-1998; 98US-0090444P.
PR 24-JUN-1998; 98US-0090445P.
PR 24-JUN-1998; 98US-0090461P.
PR 24-JUN-1998; 98US-0090472P.
PR 24-JUN-1998; 98US-0090535P.
PR 24-JUN-1998; 98US-0090538P.
PR 24-JUN-1998; 98US-0090540P.
PR 24-JUN-1998; 98US-0090557P.
PR 25-JUN-1998; 98US-0090576P.
PR 25-JUN-1998; 98US-0090578P.
PR 25-JUN-1998; 98US-0090688P.
PR 25-JUN-1998; 98US-0090690P.
PR 25-JUN-1998; 98US-0090691P.
PR 25-JUN-1998; 98US-0090694P.
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PR 25-JUN-1998; 98US-0090695P.
PR 25-JUN-1998; 98US-0090696P.
PR 26-JUN-1998; 98US-0090862P.
PR 26-JUN-1998; 98US-0090863P.
PR 01-JUL-1998; 98US-0091358P.
PR 01-JUL-1998; 98US-0091360P.
PR 02-JUL-1998; 98US-0091478P.
PR 02-JUL-1998; 98US-0091486P.
PR 02-JUL-1998; 98US-0091519P.
PR 02-JUL-1998; 98US-0091544P.
PR 02-JUL-1998; 98US-0091626P.
PR 02-JUL-1998; 98US-0091633P.
PR 02-JUL-1998; 98US-0091638P.
PR 02-JUL-1998; 98US-0091646P.
PR 02-JUL-1998; 98US-0091673P.
PR 07-JUL-1998; 98US-0091978P.
PR 07-JUL-1998; 98US-0091982P.
PR 09-JUL-1998; 98US-0092182P.
PR 10-JUL-1998; 98US-0092472P.
PR 20-JUL-1998; 98US-0093339P.
PR 30-JUL-1998; 98US-0094651P.
PR 04-AUG-1998; 98US-0095282P.
PR 04-AUG-1998; 98US-0095285P.
PR 04-AUG-1998; 98US-0095301P.
PR 04-AUG-1998; 98US-0095302P.
PR 04-AUG-1998; 98US-0095318P.
PR 04-AUG-1998; 98US-0095321P.
PR 04-AUG-1998; 98US-0095325P.
PR 10-AUG-1998; 98US-0095916P.
PR 10-AUG-1998; 98US-0095929P.
PR 10-AUG-1998; 98US-0096012P.
PR 11-AUG-1998; 98US-0096143P.
PR 11-AUG-1998; 98US-0096146P.
PR 12-AUG-1998; 98US-0096329P.
PR 17-AUG-1998; 98US-0096757P.
PR 17-AUG-1998; 98US-0096766P.
PR 17-AUG-1998; 98US-0096768P.
PR 17-AUG-1998; 98US-0096773P.
PR 17-AUG-1998; 98US-0096791P.
PR 17-AUG-1998; 98US-0096867P.
PR 17-AUG-1998; 98US-0096891P.
PR 17-AUG-1998; 98US-0096894P.
PR 17-AUG-1998; 98US-0096895P.
PR 17-AUG-1998; 98US-0096897P.
PR 18-AUG-1998; 98US-0096949P.
PR 18-AUG-1998; 98US-0096950P.
PR 18-AUG-1998; 98US-0096959P.
PR 18-AUG-1998; 98US-0096960P.
PR 18-AUG-1998; 98US-0097022P.
PR 19-AUG-1998; 98US-0097141P.
PR 20-AUG-1998; 98US-0097218P.
PR 20-AUG-1998; 98US-0097661P.
PR 26-AUG-1998; 98US-0097951P.
PR 26-AUG-1998; 98US-0097952P.
PR 26-AUG-1998; 98US-0097954P.
PR 26-AUG-1998; 98US-0097955P.
PR 26-AUG-1998; 98US-0097971P.
PR 26-AUG-1998; 98US-0097974P.
PR 26-AUG-1998; 98US-0097978P.
PR 26-AUG-1998; 98US-0097979P.
PR 26-AUG-1998; 98US-0097986P.
PR 26-AUG-1998; 98US-0098014P.
PR 31-AUG-1998; 98US-0098525P.
PR 16-SEP-1998; 98US-0100634P.
PR 12-JAN-1999; 99US-0115565P.
PA (GETH ) GENENTECH INC.
XX Baker K, Chen J, Goddard A,
XX Wood WI, Yuan J;
XX WPI; 2000-072883/06.
XX N-PSDB; AAZ64983.
XX
```

Membrane-bound proteins and related nucleotide sequences.

Claim 12; Fig 68; 822pp; English.

The invention provides membrane-bound PRO polypeptides and polynucleotides encoding them. The PRO sequences of the invention were identified based on extracellular domain homology screening. The PRO sequences have homology with proteins including LDL receptors, TIE ligands and various enzymes. The membrane-bound proteins and receptor molecules are useful as pharmaceutical and diagnostic agents. Receptor immunoadhesins, for instance, can be used as therapeutic agents to block receptor-ligand interactions. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. The PRO encoding sequences are useful as hybridization probes, in chromosome and gene mapping and in the generation of antisense RNA and DNA. PRO nucleic acid sequences will also be useful for the preparation of PRO polypeptides, especially by recombinant techniques

Sequence 123 AA;

Query Match 100.0%; Score 538; DB 3; Length 123;  
Best Local Similarity 100.0%; Pred.No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEDDHHRPADIP 60  
DB 25 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEDDHHRPADIP 84  
QY 61 DRFSAKDEAHNACVLTISPQEDDADYICSVGIGFSP 99  
DB 85 DRFSAKDEAHNACVLTISPQEDDADYICSVGIGFSP 123

RESULT 4  
AAB24061  
ID AAB24061 standard; protein; 123 AA.  
XX  
AC AAB24061;  
XX  
DT 29-JAN-2001 (first entry)  
XX  
DE Human PRO619 protein sequence SEQ ID NO:16.  
XX  
KW Human; tumour; diagnosis; neoplastic disease; neoplastic cell growth;  
KW proliferation; tumorigenesis; identification; cancer; cytostatic;  
KW neurotropic; neuroprotective; antiinflammatory; immunosuppressive;  
KW immunostimulant; antiangiogenic; leukaemia; lymphoid malignancy;  
KW neuronal disorder; glial disorder; astrocytal disorder; angiogenic;  
KW hypothalamic disorder; glandular disorder; macrophagal disorder;  
KW epithelial disorder; stromal disorder; immunologic disorder;  
XX  
OS Homo sapiens.  
XX  
PN WO2000053755-A2.  
XX  
PD 14-SEP-2000.  
XX  
PF 06-JAN-2000; 2000WO-US000376.  
XX  
PR 08-MAR-1999; 99WO-US005028.  
PR 02-JUN-1999; 99WO-US012252.  
PR 23-JUN-1999; 99US-0141037P.  
PR 07-JUL-1999; 99US-0143048P.  
PR 26-JUL-1999; 99US-0145688P.  
PR 30-NOV-1999; 99WO-US028313.  
PR 20-DEC-1999; 99WO-US030911.  
PR 05-JAN-2000; 2000WO-US0000219.  
XX  
PA (GETH ) GENENTECH INC.  
XX

PI Ashkenazi AJ, Baker KP, Goddard A, Gurney AL, Hillan KJ, Roy MA;  
PI Watanabe CK, Wood WI;  
XX WPI; 2000-572270/53.  
DR N-PSDB; AAC58371.  
XX  
XX Thirty PRO polynucleotides encoding PRO polypeptides, useful in the  
PT treatment, diagnosis and prevention of cancer.  
PT  
XX  
XX Claim 61; Fig 10; 286pp; English.  
XX  
XX The present invention describes an isolated antibody that binds to one of  
CC the human PRO proteins designated PRO212, PRO290, PRO341, PRO535, PRO619,  
CC PRO717, PRO809, PRO830, PRO848, PRO943, PRO1005, PRO1009, PRO1035,  
CC PRO1030, PRO1097, PRO1107, PRO1187, PRO1188, PRO1189, PRO1187,  
CC PRO1281, PRO23, PRO39, PRO834, PRO1317, PRO1710, PRO2094, PRO2145 OR  
CC PRO2198. PRO antagonists can be used to inhibit tumour cell growth. The  
CC PRO polypeptides and nucleotides are useful in the treatment, diagnosis  
CC and prevention of cancer. The antibodies and other anti-tumour compounds  
CC maybe used to treat various conditions, including those characterised by  
CC overexpression and/or activation of the amplified PRO genes. Exemplary  
CC conditions or disorders to be treated with such antibodies and other  
CC compounds include benign or malignant tumours (e.g., renal, liver,  
CC kidney, bladder, breast, gastric, ovarian, colorectal, prostate,  
CC pancreatic, lung, vulva, thyroid, hepatic carcinomas, sarcomas,  
CC Glioblastomas, and various head and neck tumours), leukaemias and  
CC lymphoid malignancies. Other disorders such as neuronal, glial,  
CC astrocytal, hypothalamic and other glandular, macrophagal, epithelial,  
CC stromal and blastocoele disorders, and inflammatory, angiogenic and  
CC immunologic disorders. AAC58242 to AAC58366 represent PCR primers and  
CC hybridisation probes used in the isolation of the human PRO sequences.  
CC AAC58367 to AAC58396 and AAB24057 to AAB24089 represent human PRO  
CC polynucleotide and protein sequences given in the exemplification of the  
CC present invention  
XX  
XX Sequence 123 AA;  
SQ  
Query Match 100.0%; Score 538; DB 3; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 LDALLVFPQVQALSCVLTSPQHVITRDYGVSWVQORAGSAPRYLLVYRSEEDHRRADIP 60  
Db 25 LDALLVFPQVQALSCVLTSPQHVITRDYGVSWVQORAGSAPRYLLVYRSEEDHRRADIP 84  
Qy 61 DRFSAKDEAHNACVLIISVPQEDDADYVCSVGYGFSP 99  
Db 85 DRFSAKDEAHNACVLIISVPQEDDADYVCSVGYGFSP 123  
RESULT 5  
AAU12372  
ID AAU12372 standard; protein; 123 AA.  
XX  
XX AAU12372;  
XX  
XX 24-OCT-2001 (first entry)  
XX  
XX Human PRO619 polypeptide sequence.  
XX  
XX Human secretory and transmembrane; PRO; mammalian; cancer; lung; breast;  
KW prostate; cervical; tumour necrosis factor-alpha; TNF-alpha; cartilage;  
KW ear; proliferation; glucose; free fatty acid; skeletal muscle; adipocyte;  
KW A-peptide; factor VIIA; gene therapy.  
XX  
XX Homo sapiens.  
OS  
XX  
XX WO2000140466-A2.  
XX  
XX 07-JUN-2001.  
PD  
XX  
XX 01-DEC-2000; 2000WO-US032678.  
PF  
XX

PR 01-DEC-1999; 99WO-US028301.  
PR 01-DEC-1999; 99WO-US028634.  
PR 02-DEC-1999; 99WO-US028551.  
PR 02-DEC-1999; 99WO-US028564.  
PR 09-DEC-1999; 99US-0170262P.  
PR 16-DEC-1999; 99WO-US030095.  
PR 20-DEC-1999; 99WO-US030311.  
PR 20-DEC-1999; 99WO-US030399.  
PR 30-DEC-1999; 99WO-US031443.  
PR 30-DEC-1999; 99WO-US031274.  
PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000277.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 18-FEB-2000; 2000WO-US004341.  
PR 18-FEB-2000; 2000WO-US004342.  
PR 22-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 24-FEB-2000; 2000WO-US005004.  
PR 01-MAR-2000; 2000WO-US005601.  
PR 02-MAR-2000; 2000WO-US005841.  
PR 03-MAR-2000; 2000US-0187202P.  
PR 10-MAR-2000; 2000WO-US006319.  
PR 15-MAR-2000; 2000WO-US006884.  
PR 20-MAR-2000; 2000WO-US007377.  
PR 21-MAR-2000; 2000WO-US007532.  
PR 30-MAR-2000; 2000WO-US008439.  
PR 17-MAY-2000; 2000WO-US013705.  
PR 22-MAY-2000; 2000WO-US014042.  
PR 30-MAY-2000; 2000WO-US014941.  
PR 02-JUN-2000; 2000US-0209832P.  
PR 05-JUN-2000; 2000US-0209832P.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US022031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023528.  
PR 08-NOV-2000; 2000WO-US030952.  
PR 10-NOV-2000; 2000WO-US030873.  
XX  
XX (GETH ) GENENTECH INC.  
XX  
XX Baker KP, Beresini M, DeForge L, Desnoyers L, Filvaroff E, Gao W;  
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;  
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;  
XX  
XX WPI; 2001-408281/43.  
DR N-PSDB; AAS21444.  
XX  
XX Isolated , secretory and transmembrane PRO polypeptide used to detect  
PT other PRO polypeptides, link bioactive molecules to cells expressing PRO  
PT polypeptides, and detect the presence of mammalian tumors e.g. lung,  
PT breast, prostate, cervical.  
XX  
XX Claim 12; Fig 402; 813pp; English.  
XX  
XX AAU12172-AAU12446 represent novel human secretory and transmembrane PRO  
CC polypeptides. The PRO polypeptides are useful to detect other PRO  
CC polypeptides, to link bioactive molecules to cells expressing PRO  
CC polypeptides, to modulate biological activities of cells expressing PRO  
CC polypeptides, and to detect the presence of mammalian lung, colon,  
CC breast, prostate, rectal, cervical or liver tumours by comparing PRO  
CC polypeptide expression in a cell sample to that in a control sample. Some  
CC of the 275 sequences are also useful to stimulate the release of tumour  
CC necrosis factor-alpha (TNF-alpha) from human blood, the proliferation or  
CC differentiation of chondrocytes, the proliferation or gene expression in  
CC pericyte cells, the release of proteoglycans from cartilage, the  
CC proliferation of inner ear utricular supporting cells or of T-  
CC lymphocytes, the release of a cytokine from peripheral blood monocytes  
CC (PBMCs), or the proliferation of endothelial cells. Some of the PRO  
CC polypeptides may modulate glucose or free fatty acid uptake by skeletal  
CC muscle cells or by adipocytes; or inhibit binding of A-peptide to factor  
CC VIIA. The PRO polypeptides can be used in assays to identify molecules

CC involved in binding interactions. The polynucleotides encoding PRO  
CC polypeptides can be used to generate probes, antisense RNA/DNA,  
CC transgenic or knock out animals and can be used in gene therapy  
XX  
SQ Sequence 123 AA;  
Query Match 100.0%; Score 538; DB 4; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEDHRRPADIP 60  
DB 25 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEDHRRPADIP 84  
QY 61 DRFSAKDEAHNACVLTISVPQEDDADYCSVGYGFSP 99  
DB 85 DRFSAKDEAHNACVLTISVPQEDDADYCSVGYGFSP 123  
RESULT 6  
AAB65178  
ID AAB65178 standard; protein; 123 AA.  
AC AAB65178;  
DT 02-APR-2001 (first entry)  
XX  
XX  
DE Human PRO619 (UNQ355) protein sequence SEQ ID NO:117.  
KW Human; secreted and transmembrane protein; PRO; cytostatic; cell death;  
KW cancer; chromosomal mapping; gene mapping; tissue typing;  
KW diagnostic assay.  
XX  
XX Homo sapiens.  
XX WO200073454-A1.  
XX 07-DEC-2000.  
XX  
XX 30-MAR-2000; 2000WO-US008439.  
XX  
PR 02-JUN-1999; 99WO-US012252.  
PR 23-JUN-1999; 99US-0141037P.  
PR 07-JUL-1999; 99US-0143048P.  
PR 20-JUL-1999; 99US-0144758P.  
PR 26-JUL-1999; 99US-0145698P.  
PR 28-JUL-1999; 99US-0146222P.  
PR 17-AUG-1999; 99US-0149396P.  
PR 15-SEP-1999; 99WO-US021090.  
PR 15-SEP-1999; 99WO-US021547.  
PR 08-OCT-1999; 99US-0158663P.  
PR 30-NOV-1999; 99WO-US028313.  
PR 01-DEC-1999; 99WO-US028301.  
PR 16-DEC-1999; 99WO-US030095.  
PR 20-DEC-1999; 99WO-US030911.  
PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 18-FEB-2000; 2000WO-US004341.  
PR 22-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 24-FEB-2000; 2000WO-US005004.  
PR 02-MAR-2000; 2000WO-US005541.  
PR 15-MAR-2000; 2000WO-US006584.  
PR 20-MAR-2000; 2000WO-US007377.  
XX  
XX (GETH ) GENENTECH INC.  
XX  
XX Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;  
XX Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;  
XX Grimaldi CJ, Gurney AL, Kijavini IJ, Napier MA, Pan J, Paoni NF;  
XX Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;  
XX Zhang Z;

XX WPI: 2001-032160/04.  
DR N-PSDB; AAF44129.  
XX  
PT PRO polynucleotides used to produce polypeptides used to target bioactive  
PT molecules such as toxins, radiolabels or antibodies, to specific cells,  
PT to cause targeted cell death.  
XX  
PS Claim 12; Fig 68; 935pp; English.  
XX  
CC The present invention describes human secreted and transmembrane PRO  
CC proteins. The PRO proteins have cytostatic activity. The PRO proteins can  
CC be used for targeted delivery of bioactive molecules, such as toxins,  
CC radiolabels or antibodies, that cause cell death. PRO nucleotide  
CC sequences, and their fragments, can be used as hybridisation probes, in  
CC chromosomal and gene mapping, and in the generation of anti-sense RNA and  
CC DNA. They may also be used to produce transgenic animals which are used  
CC to develop and screen therapeutically useful reagents. The PRO nucleotide  
CC and protein sequence can be used for tissue typing and in treating  
CC cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to  
CC AAF44470 represent PCR primers and hybridisation probes used in the  
CC isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65154 to  
CC AAB65300 represent human PRO polynucleotide and protein sequences given  
CC in the exemplification of the present invention  
XX  
SQ Sequence 123 AA;  
Query Match 100.0%; Score 538; DB 4; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEDHRRPADIP 60  
DB 25 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEDHRRPADIP 84  
QY 61 DRFSAKDEAHNACVLTISVPQEDDADYCSVGYGFSP 99  
DB 85 DRFSAKDEAHNACVLTISVPQEDDADYCSVGYGFSP 123  
RESULT 7  
AAB65178  
ID AAB65178 standard; protein; 123 AA.  
AC AAB65178;  
DT 14-APR-2003 (first entry)  
XX  
XX Human PRO polypeptide #25.  
XX  
XX Human; PRO; cytostatic; tumour; cancer; breast; lung; stomach; liver;  
XX horse; cow; dog; cat; sheep; pig; goat; rabbit; ADEPT;  
XX antibody-dependent enzyme mediated prodrug therapy.  
XX  
XX Homo sapiens.  
XX US2003027163-A1.  
XX  
XX 06-FEB-2003.  
XX  
XX 15-NOV-2001; 2001US-00997666.  
XX  
XX 16-JUN-1997; 97US-0049787P.  
XX 17-OCT-1997; 97US-0062250P.  
XX 05-NOV-1997; 97WO-US020069.  
XX 12-NOV-1997; 97US-0055186P.  
XX 13-NOV-1997; 97US-0065311P.  
XX 24-NOV-1997; 97US-0066770P.  
XX 25-FEB-1998; 98US-0075945P.  
XX 20-MAR-1998; 98US-0078910P.  
XX 28-APR-1998; 98US-0083322P.  
XX 07-MAY-1998; 98US-0084600P.  
XX 28-MAY-1998; 98US-0087106P.







PR 02-JUN-2000; 2000WO-US015264.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US022031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023328.  
PR 01-DEC-2000; 2000WO-US030952.  
PR 08-NOV-2000; 2000WO-US032678.  
PR 28-FEB-2001; 2001WO-US006520.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 20-JUN-2001; 2001WO-US019892.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 28-AUG-2001; 2001US-00941992.  
XX XX  
PA (GETH ) GENENTECH INC.  
XX XX  
PI Askenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;  
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;  
PI Grimaldi JC, Gurney AU, Klabavin IU, Napier MA, Pan J, Paoni NF;  
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;  
PI Zhang Z;  
XX XX  
XX WPI; 2003-247083/24.  
DR N-PSDB; ABX80196.  
XX XX  
PT Novel isolated PRO polypeptides e.g., PRO826, PRO1068, PRO1184, PRO1346  
PT and PRO1375, which stimulate proliferation of stimulated T-lymphocytes  
PT are therapeutically useful for enhancing immune response and in cancer  
PT treatments.  
XX XX  
FS Claim 12; Fig 68; 648pp; English.  
XX XX  
CC The invention describes an isolated human PRO polypeptide. The PRO  
CC polypeptides are useful in detecting PRO polypeptides in a sample, in  
CC linking a bioactive molecule to a cell expressing a PRO polypeptide, and  
CC in modulating at least one biological activity of a cell expressing a PRO  
CC polypeptide. PRO1312 stimulates hypertrophy of neonatal heart and is thus  
CC useful for treating cardiac insufficiency disorders. PRO1154 and PRO1186  
CC stimulate adrenal cortical capillary endothelial growth, and PRO336,  
CC PRO943, PRO828, PRO826, PRO1068 or PRO335, PRO826, PRO819, PRO1126,  
CC PRO1360 and PRO1387 induce c-fos in endothelial cells, and are thus  
CC useful for treating conditions or disorders where angiogenesis would be  
CC beneficial, e.g. wound healing and antagonist of this polypeptide are  
CC useful for treating cancerous tumours. PRO812 inhibits vascular  
CC endothelial growth factor (VEGF) stimulated proliferation of endothelial  
CC cells and is thus useful for inhibiting endothelial cell growth in  
CC mammals which would be beneficial in inhibiting tumour growth. PRO826,  
CC PRO1068, PRO1184, PRO1346 and PRO1375 stimulate proliferation of  
CC stimulated T-lymphocytes and are therapeutically useful for enhancing  
CC immune response. PRO828, PRO826, PRO1068 or PRO132 enhance survival of  
CC retinal neurons cells (PRO1132 is also enhances survival/proliferation of  
CC rod photoreceptor cells) and therefore are useful for treating retinal  
CC disorders of injuries, e.g. retinitis pigmentosa, AMD. PRO819, PRO813  
CC and PRO1066 induce proliferation of mammalian kidney mesangial cells,  
CC and therefore are useful for treating kidney disorders associated with  
CC decreased mesangial cell function such as Berger disease or other  
CC nephropathies associated with dermatitis, herpeticiformis or Crohn's  
CC disease. PRO1310, PRO844, PRO1311, PRO1192 and PRO1387 induce the  
CC proliferation and/or redifferentiation of chondrocytes in culture and are  
CC thus useful for treating sports injuries, and arthritis. This is the  
CC amino acid sequence of a novel human PRO protein  
XX XX  
SQ Sequence 123 AA;  
Query Match 100.0%; Score 538; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEDHHRPADIP 60  
DB 25 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEDHHRPADIP 84  
OY 61 DRFSAKDEAHNACVLTISPQEDDADYCYCSGVGFSP 99

Db 85 DRFSAKDEAHNACVLTISPQEDDADYCYCSGVGFSP 123  
|||||  
RESULT 9  
ABU82583  
ID ABU82583 standard; protein; 123 AA.  
XX XX  
AC ABU82583;  
XX XX  
DT 26-JUN-2003 (first entry)  
XX XX  
DE Human secreted/transmembrane protein PRO619.  
XX XX  
KW Human; PRO; secreted protein; transmembrane protein;  
KW cardiac insufficiency disorders; angiogenesis; wound healing;  
KW cancerous tumour; immune response; retinal disorder; sight loss;  
KW retinitis pigmentosa; age-related macular degeneration; AMD;  
KW kidney disorder; Berger disease; nephropathy; dermatitis; herpeticiformis;  
KW Crohn's disease; sports injury; arthritis.  
XX OS  
XX Homo sapiens.  
XX PN  
XX US2003032023-A1.  
XX XX  
PD 13-FEB-2003.  
XX PF  
PF 14-NOV-2001; 2001US-00990711.  
XX XX  
PR 16-JUN-1997; 97US-0049787P.  
PR 17-OCT-1997; 97US-0062250P.  
PR 05-NOV-1997; 97WO-US020069.  
PR 12-NOV-1997; 97US-0065186P.  
PR 13-NOV-1997; 97US-0065311P.  
PR 24-NOV-1997; 97US-0066770P.  
PR 25-FEB-1998; 98US-0075945P.  
PR 20-MAR-1998; 98US-0078910P.  
PR 28-APR-1998; 98US-0083322P.  
PR 07-MAY-1998; 98US-0084600P.  
PR 28-MAY-1998; 98US-0087106P.  
PR 02-JUN-1998; 98US-0087607P.  
PR 02-JUN-1998; 98US-0087609P.  
PR 02-JUN-1998; 98US-0087759P.  
PR 03-JUN-1998; 98US-0087827P.  
PR 04-JUN-1998; 98US-0088021P.  
PR 04-JUN-1998; 98US-0088025P.  
PR 04-JUN-1998; 98US-0088026P.  
PR 04-JUN-1998; 98US-0088028P.  
PR 04-JUN-1998; 98US-0088029P.  
PR 04-JUN-1998; 98US-0088030P.  
PR 04-JUN-1998; 98US-0088033P.  
PR 04-JUN-1998; 98US-0088326P.  
PR 05-JUN-1998; 98US-0088167P.  
PR 05-JUN-1998; 98US-0088202P.  
PR 05-JUN-1998; 98US-0088212P.  
PR 05-JUN-1998; 98US-0088217P.  
PR 09-JUN-1998; 98US-0088655P.  
PR 10-JUN-1998; 98US-0088734P.  
PR 10-JUN-1998; 98US-0088738P.  
PR 10-JUN-1998; 98US-0088742P.  
PR 10-JUN-1998; 98US-0088810P.  
PR 10-JUN-1998; 98US-0088824P.  
PR 10-JUN-1998; 98US-0088826P.  
PR 11-JUN-1998; 98US-0088858P.  
PR 11-JUN-1998; 98US-0088861P.  
PR 11-JUN-1998; 98US-0088876P.  
PR 12-JUN-1998; 98US-0089105P.  
PR 16-JUN-1998; 98US-0089440P.  
PR 16-JUN-1998; 98US-0089512P.  
PR 16-JUN-1998; 98US-0089514P.  
PR 17-JUN-1998; 98US-0089532P.  
PR 17-JUN-1998; 98US-0089538P.  
PR 17-JUN-1998; 98US-0089598P.



RESULT 10  
 ID ABO17816 standard; protein; 123 AA.  
 AC ABO17816;  
 XX  
 DT 26-AUG-2003 (first entry)  
 XX  
 DE Novel human secreted and transmembrane protein PRO619.  
 XX  
 KW Human; secreted and transmembrane protein; PRO; antiinflammatory;  
 KW antiarteriosclerotic; cardiant; anti-infertility; anti-HIV; cytostatic;  
 KW antidiabetic; gene therapy; tumour necrosis factor (TNF)-alpha release;  
 KW TNF-alpha release; cell proliferation; cell differentiation;  
 KW Gene expression modulator; proteoglycan release; cytokine release;  
 KW tumour; inflammatory disease; organ failure; atherosclerosis;  
 KW cardiac injury; infertility; birth defect; premature aging; AIDS;  
 KW acquired immunodeficiency syndrome; cancer; diabetic complication;  
 KW chromosome mapping; gene mapping; pharmaceutical; diagnostic; biosensor;  
 KW bioreactor; tissue typing.  
 XX  
 OS Homo sapiens.  
 XX  
 XX US2003032156-A1.  
 XX  
 XX 13-FEB-2003.  
 XX  
 XX 06-MAY-2002; 2002US-00140474.  
 XX  
 31-MAR-1997; 97WO-US005230.  
 PR 12-JUN-1998; 98WO-US012456.  
 PR 14-JUL-1998; 98WO-US014552.  
 PR 28-AUG-1998; 98WO-US017888.  
 PR 10-SEP-1998; 98WO-US018824.  
 PR 14-SEP-1998; 98WO-US019094.  
 PR 14-SEP-1998; 98WO-US019177.  
 PR 16-SEP-1998; 98WO-US019330.  
 PR 17-SEP-1998; 98WO-US019437.  
 PR 07-OCT-1998; 98WO-US021141.  
 PR 29-OCT-1998; 98WO-US022591.  
 PR 29-OCT-1998; 98WO-US022592.  
 PR 20-NOV-1998; 98WO-US024855.  
 PR 01-DEC-1998; 98WO-US025108.  
 PR 05-JAN-1999; 99WO-US000106.  
 PR 08-MAR-1999; 99WO-US005028.  
 PR 10-MAR-1999; 99WO-US005190.  
 PR 20-APR-1999; 99WO-US008615.  
 PR 14-MAY-1999; 99WO-US010733.  
 PR 02-JUN-1999; 99WO-US012252.  
 PR 01-SEP-1999; 99WO-US020111.  
 PR 08-SEP-1999; 99WO-US020594.  
 PR 13-SEP-1999; 99WO-US020944.  
 PR 15-SEP-1999; 99WO-US021090.  
 PR 15-SEP-1999; 99WO-US021547.  
 PR 05-OCT-1999; 99WO-US023089.  
 PR 29-NOV-1999; 99WO-US028214.  
 PR 30-NOV-1999; 99WO-US028313.  
 PR 30-NOV-1999; 99WO-US028409.  
 PR 01-DEC-1999; 99WO-US028301.  
 PR 01-DEC-1999; 99WO-US028634.  
 PR 02-DEC-1999; 99WO-US028551.  
 PR 02-DEC-1999; 99WO-US028564.  
 PR 02-DEC-1999; 99WO-US028565.  
 PR 16-DEC-1999; 99WO-US030095.  
 PR 20-DEC-1999; 99WO-US030911.  
 PR 20-DEC-1999; 99WO-US030999.  
 PR 22-DEC-1999; 99WO-US030720.  
 PR 30-DEC-1999; 99WO-US031243.  
 PR 30-DEC-1999; 99WO-US031274.  
 PR 05-JAN-2000; 2000WO-US000219.  
 PR 06-JAN-2000; 2000WO-US000277.  
 PR 06-JAN-2000; 2000WO-US000376.  
 PR 11-FEB-2000; 2000WO-US003565.  
 PR 18-FEB-2000; 2000WO-US004341.  
 PR 18-FEB-2000; 2000WO-US004342.  
 PR 22-FEB-2000; 2000WO-US004414.  
 PR 24-FEB-2000; 2000WO-US004914.  
 PR 24-FEB-2000; 2000WO-US005004.  
 PR 01-MAR-2000; 2000WO-US005601.  
 PR 02-MAR-2000; 2000WO-US005746.  
 PR 02-MAR-2000; 2000WO-US005841.  
 PR 10-MAR-2000; 2000WO-US006319.  
 PR 15-MAR-2000; 2000WO-US006884.  
 PR 20-MAR-2000; 2000WO-US007377.  
 PR 21-MAR-2000; 2000WO-US007532.  
 PR 30-MAR-2000; 2000WO-US008439.  
 PR 17-MAY-2000; 2000WO-US013705.  
 PR 22-MAY-2000; 2000WO-US014042.  
 PR 30-MAY-2000; 2000WO-US014941.  
 PR 02-JUN-2000; 2000WO-US015264.  
 PR 28-JUL-2000; 2000WO-US020710.  
 PR 11-AUG-2000; 2000WO-US020231.  
 PR 23-AUG-2000; 2000WO-US023522.  
 PR 24-AUG-2000; 2000WO-US023328.  
 PR 08-NOV-2000; 2000WO-US030952.  
 PR 10-NOV-2000; 2000WO-US030873.  
 PR 01-DEC-2000; 2000WO-US032678.  
 PR 20-DEC-2000; 2000US-00747259.  
 PR 20-DEC-2000; 2000WO-US034956.  
 PR 28-FEB-2001; 2001US-00796498.  
 PR 28-FEB-2001; 2001WO-US006520.  
 PR 01-MAR-2001; 2001WO-US006666.  
 PR 09-MAR-2001; 2001US-00802706.  
 PR 14-MAR-2001; 2001US-00805689.  
 PR 22-MAR-2001; 2001US-00816744.  
 PR 05-APR-2001; 2001US-00828366.  
 PR 10-MAY-2001; 2001US-00854208.  
 PR 10-MAY-2001; 2001US-00854208.  
 PR 18-MAY-2001; 2001US-00860216.  
 PR 25-MAY-2001; 2001US-00866028.  
 PR 25-MAY-2001; 2001US-00866034.  
 PR 25-MAY-2001; 2001WO-US017092.  
 PR 01-JUN-2001; 2001US-00872035.  
 PR 01-JUN-2001; 2001WO-US017800.  
 PR 05-JUN-2001; 2001US-00874503.  
 PR 14-JUN-2001; 2001US-00882636.  
 PR 19-JUN-2001; 2001US-00886342.  
 PR 20-JUN-2001; 2001WO-US019692.  
 PR 21-JUN-2001; 2001US-00887879.  
 PR 22-JUN-2001; 2001WO-US020116.  
 PR 29-JUN-2001; 2001WO-US021066.  
 PR 09-JUL-2001; 2001WO-US021735.  
 PR 18-JUL-2001; 2001US-00908827.  
 PR 06-AUG-2001; 2001US-00924419.  
 PR 09-AUG-2001; 2001US-00927796.  
 PR 16-AUG-2001; 2001US-00931836.  
 PR 19-DEC-2001; 2001US-00028072.  
 XX  
 PA (GETH ) GENENTECH INC.  
 XX  
 XX Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;  
 PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;  
 PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;  
 XX  
 DR WPI; 2003-341980/32.  
 DR N-PSDB; ACD24053.  
 XX  
 XX New secreted and transmembrane PRO nucleic acids, for treating  
 PT inflammation, organ failure, atherosclerosis, cardiac injury,  
 PT infertility, birth defects, premature aging, acquired immunodeficiency  
 PT syndrome (AIDS), or cancer.  
 XX  
 PS Claim 12; Fig 402; 660pp; English.  
 XX

CC The invention describes an isolated nucleic acid (I) comprising, or which  
CC has 80 % sequence identity to, or the full-length coding sequence of, one  
CC of 275 nucleotide sequences, and which encodes a corresponding  
CC polypeptide selected from 275 amino acid sequences, where all sequences  
CC are given in the specification. The polypeptide encoded by (I) is used to  
CC detect PRO polypeptides, link a bioactive molecule to a cell expressing a  
CC PRO polypeptide, modulate a biological activity of a cell, stimulate the  
CC release of tumour necrosis factor (TNF)-alpha from human blood, modulate  
CC the uptake of glucose or free fatty acid by cells, stimulate or inhibit  
CC the proliferation or differentiation of cells or gene expression, or  
CC stimulate the release of proteoglycans, stimulate the release of cytokine  
CC from peripheral blood mononuclear cells, inhibit the binding of A-peptide  
CC to factor VIIA, or detect the presence of tumour in a mammal. The nucleic  
CC acid and polypeptide encoded by it, are useful for treating inflammatory  
CC diseases, organ failure, atherosclerosis, cardiac injury, infertility,  
CC birth defects, premature aging, acquired immunodeficiency syndrome  
CC (AIDS), cancer, or diabetic complications. The nucleic acid is useful as  
CC hybridisation probes, in chromosome and gene mapping, and in generating  
CC antisense RNA or DNA. The polypeptides are useful as pharmaceuticals,  
CC diagnostics, biosensors or bioreactors. Both are useful in tissue typing.  
CC This is the amino acid sequence of a novel human secreted and  
CC transmembrane PRO polypeptide

SQ Sequence 123 AA;

Query Match 100.0%; Score 538; DB 6; Length 123;

Best Local Similarity 100.0%; Pred. No. 1e-51;

Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LDALLVFPQVAQLSCLTSPQVTRIDYGVSWYQQRAGSAPRYLLVYRSEEDHRRADIP 60

DB 25 LDALLVFPQVAQLSCLTSPQVTRIDYGVSWYQQRAGSAPRYLLVYRSEEDHRRADIP 84

QY 61 DRFSAKDEAHNACVLTSIPQVEDDADYCVSGYGFSP 99

DB 85 DRFSAKDEAHNACVLTSIPQVEDDADYCVSGYGFSP 123

RESULT 11

ID ABU60502 standard; protein; 123 AA.

XX AC ABU60502;

XX DT 01-MAY-2003 (first entry)

XX DE Human secreted/transmembrane protein, #43.

XX KW Human; PRO; secreted; transmembrane; signal peptide; pharmaceutical;  
XX diagnostic; therapeutic; gene therapy.

XX OS Homo sapiens.

XX PN US2002160384-A1.

XX PD 31-OCT-2002.

XX PF 14-NOV-2001; 2001US-00992598.

XX PR 16-JUN-1997; 97US-0049787P.

XX PR 17-OCT-1997; 97US-0062250P.

XX PR 05-NOV-1997; 97WO-US020069.

XX PR 12-NOV-1997; 97US-0065186P.

XX PR 13-NOV-1997; 97US-0065311P.

XX PR 24-NOV-1997; 97US-0066770P.

XX PR 25-FEB-1998; 98US-0075945P.

XX PR 20-MAR-1998; 98US-0078910P.

XX PR 28-APR-1998; 98US-0083332P.

XX PR 07-MAY-1998; 98US-0084600P.

XX PR 28-MAY-1998; 98US-0087106P.

XX PR 02-JUN-1998; 98US-0087607P.

XX PR 02-JUN-1998; 98US-0087609P.

XX PR 02-JUN-1998; 98US-0087759P.

PR 03-JUN-1998; 98US-0087827P.  
PR 04-JUN-1998; 98US-0088021P.  
PR 04-JUN-1998; 98US-0088025P.  
PR 04-JUN-1998; 98US-0088028P.  
PR 04-JUN-1998; 98US-0088028P.  
PR 04-JUN-1998; 98US-0088029P.  
PR 04-JUN-1998; 98US-0088030P.  
PR 04-JUN-1998; 98US-0088033P.  
PR 04-JUN-1998; 98US-0088326P.  
PR 05-JUN-1998; 98US-0088167P.  
PR 05-JUN-1998; 98US-0088202P.  
PR 05-JUN-1998; 98US-0088212P.  
PR 05-JUN-1998; 98US-0088217P.  
PR 09-JUN-1998; 98US-0088655P.  
PR 10-JUN-1998; 98US-0088734P.  
PR 10-JUN-1998; 98US-0088738P.  
PR 10-JUN-1998; 98US-0088743P.  
PR 10-JUN-1998; 98US-0088810P.  
PR 10-JUN-1998; 98US-0088824P.  
PR 10-JUN-1998; 98US-0088826P.  
PR 11-JUN-1998; 98US-0088858P.  
PR 11-JUN-1998; 98US-0088861P.  
PR 11-JUN-1998; 98US-0088876P.  
PR 12-JUN-1998; 98US-0089105P.  
PR 12-JUN-1998; 98US-0089440P.  
PR 16-JUN-1998; 98US-0089512P.  
PR 16-JUN-1998; 98US-0089514P.  
PR 17-JUN-1998; 98US-0089532P.  
PR 17-JUN-1998; 98US-0089538P.  
PR 17-JUN-1998; 98US-0089598P.  
PR 17-JUN-1998; 98US-0089599P.  
PR 17-JUN-1998; 98US-0089600P.  
PR 17-JUN-1998; 98US-0089653P.  
PR 18-JUN-1998; 98US-0089801P.  
PR 18-JUN-1998; 98US-0089907P.  
PR 18-JUN-1998; 98US-0089908P.  
PR 16-SEP-1998; 98WO-US019330.  
PR 17-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 01-DEC-1998; 98WO-US025108.  
PR 05-JAN-1999; 99WO-US000106.  
PR 08-MAR-1999; 99WO-US005028.  
PR 02-JUN-1999; 99WO-US012252.  
PR 15-SEP-1999; 99WO-US021090.  
PR 15-SEP-1999; 99WO-US021547.  
PR 30-NOV-1999; 99WO-US028313.  
PR 01-DEC-1999; 99WO-US028301.  
PR 01-DEC-1999; 99WO-US028634.  
PR 16-DEC-1999; 99WO-US030095.  
PR 20-DEC-1999; 99WO-US030911.  
PR 05-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 11-FEB-2000; 2000WO-US004341.  
PR 22-FEB-2000; 2000WO-US004414.  
PR 24-FEB-2000; 2000WO-US004914.  
PR 02-MAR-2000; 2000WO-US005004.  
PR 10-MAR-2000; 2000WO-US005841.  
PR 15-MAR-2000; 2000WO-US006884.  
PR 20-MAR-2000; 2000WO-US007377.  
PR 30-MAR-2000; 2000WO-US008439.  
PR 15-MAY-2000; 2000WO-US013358.  
PR 17-MAY-2000; 2000WO-US013705.  
PR 22-MAY-2000; 2000WO-US014042.  
PR 30-MAY-2000; 2000WO-US014941.  
PR 02-JUN-2000; 2000WO-US015264.  
PR 28-JUL-2000; 2000WO-US020710.  
PR 11-AUG-2000; 2000WO-US022031.  
PR 23-AUG-2000; 2000WO-US023522.  
PR 24-AUG-2000; 2000WO-US023328.  
PR 08-NOV-2000; 2000WO-US030952.  
PR 01-DEC-2000; 2000WO-US032678.

PR 28-FEB-2001; 2001WO-US006520.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 20-JUN-2001; 2001WO-US019692.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 28-AUG-2001; 2001US-00941992.  
XX XX  
PA (GETH ) GENENTECH INC.  
XX XX  
PI Ashkenazi AJ, Baker KP, Botstein D, Desnovers L, Eaton DL,  
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ,  
PI Grimaldi JC, Garney AL, Kijavini LJ, Napier MA, Pan J, Paoni NF,  
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI,  
PI Zhang Z;  
XX XX  
DR WPI; 2003-288106/28.  
DR N-PSDB; ABX90174.  
XX XX  
PT New transmembrane polypeptides and nucleic acids encoding the  
PT polypeptides, useful in gene therapy, in chromosome identification, as  
PT chromosome markers, or in generating probes.  
XX XX  
PS Claim 12; Fig 68; 650pp; English.  
XX XX  
CC The invention discloses isolated PRO secreted/transmembrane polypeptides  
CC comprising a sequence without signal peptide and the nucleic acid  
CC encoding them. The polypeptides can be used to raise antibodies that  
CC specifically bind to the PRO polypeptide, for linking a bioactive  
CC molecule to a cell expressing a PRO protein and for modulating at least  
CC one biological activity of a cell. The PRO polypeptides or  
CC polynucleotides are also useful in gene therapy, in chromosome  
CC identification, as chromosome markers, or in generating probes. The PRO  
CC polypeptides are useful as molecular markers for protein electrophoresis,  
CC and the isolated nucleic acids may be used for recombinantly expressing  
CC those markers. The PRO polypeptides and nucleic acids may also be used in  
CC tissue typing. Anti-PRO antibodies are useful in diagnostic assays for  
CC PRO, and in affinity purification of PRO from recombinant cell culture or  
CC natural sources. The sequences presented in ASU60478-ABU60624 are the PRO  
CC polynucleotides of the invention. Note: The sequence data for this patent  
CC is also available in electronic format from USFTO at  
CC seqdata.uspto.gov/sequence.html  
XX XX  
SQ Sequence 123 AA;  
  
Query Match 100.0%; Score 538; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Caps 0;  
  
Qy 1 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEEDHRRPADIP 60  
Db 25 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEEDHRRPADIP 84  
  
Qy 61 DRFSAAXDEAHNACVLTISPVPQEDDADYCVSGYGFSF 99  
Db 85 DRFSAAXDEAHNACVLTISPVPQEDDADYCVSGYGFSF 123  
  
RESULT 12  
ABU13884  
ID ABU13884 standard; protein; 123 AA.  
XX XX  
AC ABU13884;  
XX XX  
DT 26-FEB-2003 (first entry)  
DE Human PRO619 polypeptide.  
XX XX  
KW Human; PRO polypeptide; secreted protein; transmembrane protein;  
KW genetic disorder; antibacterial; immunosuppressive.  
XX XX  
OS Homo sapiens.  
XX XX  
PN US2002103125-A1.

XX PD  
XX PF  
XX PF  
XX XX  
PR 01-AUG-2002.  
PR 20-NOV-2001; 2001US-00989731.  
PR 16-JUN-1997; 97US-0049787P.  
PR 17-OCT-1997; 97US-0062250P.  
PR 05-NOV-1997; 97WO-US020069.  
PR 12-NOV-1997; 97US-0065186P.  
PR 13-NOV-1997; 97US-0065311P.  
PR 24-NOV-1997; 97US-0066770P.  
PR 25-FEB-1998; 98US-0075945P.  
PR 20-MAR-1998; 98US-0078910P.  
PR 28-APR-1998; 98US-0083322P.  
PR 07-MAY-1998; 98US-0084600P.  
PR 28-MAY-1998; 98US-0087106P.  
PR 02-JUN-1998; 98US-0087609P.  
PR 02-JUN-1998; 98US-0087759P.  
PR 03-JUN-1998; 98US-0087827P.  
PR 04-JUN-1998; 98US-0088021P.  
PR 04-JUN-1998; 98US-0088025P.  
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PR 04-JUN-1998; 98US-0088035P.  
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PR 10-JUN-1998; 98US-0088738P.  
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PR 12-JUN-1998; 98US-0089105P.  
PR 16-JUN-1998; 98US-0089440P.  
PR 16-JUN-1998; 98US-0089512P.  
PR 16-JUN-1998; 98US-0089514P.  
PR 17-JUN-1998; 98US-0089532P.  
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PR 17-JUN-1998; 98US-0089653P.  
PR 18-JUN-1998; 98US-0089801P.  
PR 18-JUN-1998; 98US-0089907P.  
PR 18-JUN-1998; 98US-0089908P.  
PR 16-SEP-1998; 98WO-US019330.  
PR 17-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 01-DEC-1998; 98WO-US025108.  
PR 05-JAN-1999; 98WO-US000106.  
PR 08-MAR-1999; 98WO-US005028.  
PR 02-JUN-1999; 98WO-US012252.  
PR 15-SEP-1999; 98WO-US021090.  
PR 15-SEP-1999; 98WO-US021547.  
PR 30-NOV-1999; 98WO-US028313.  
PR 01-DEC-1999; 98WO-US028301.  
PR 01-DEC-1999; 98WO-US028634.  
PR 16-DEC-1999; 98WO-US030095.  
PR 20-DEC-1999; 98WO-US030911.  
PR 06-JAN-2000; 2000WO-US000219.  
PR 06-JAN-2000; 2000WO-US000376.  
PR 11-FEB-2000; 2000WO-US003565.  
PR 18-FEB-2000; 2000WO-US004341.

PR 22-FEB-2000; 200OWO-US004414.  
PR 24-FEB-2000; 200OWO-US004914.  
PR 24-FEB-2000; 200OWO-US005004.  
PR 02-MAR-2000; 200OWO-US005841.  
PR 10-MAR-2000; 200OWO-US006319.  
PR 15-MAR-2000; 200OWO-US006984.  
PR 20-MAR-2000; 200OWO-US007377.  
PR 30-MAR-2000; 200OWO-US008439.  
PR 15-MAY-2000; 200OWO-US013358.  
PR 17-MAY-2000; 200OWO-US013705.  
PR 22-MAY-2000; 200OWO-US014042.  
PR 30-MAY-2000; 200OWO-US014941.  
PR 02-JUN-2000; 200OWO-US015264.  
PR 28-JUL-2000; 200OWO-US020710.  
PR 11-AUG-2000; 200OWO-US022031.  
PR 23-AUG-2000; 200OWO-US023522.  
PR 24-AUG-2000; 200OWO-US023328.  
PR 08-NOV-2000; 200OWO-US030952.  
PR 01-DEC-2000; 200OWO-US032678.  
PR 28-FEB-2001; 2001WO-US006520.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 20-JUN-2001; 2001WO-US019892.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 28-AUG-2001; 2001US-00941992.  
XX  
XX (GETH ) GENENTECH LTD.  
XX  
PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;  
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;  
PI Grimaldi JC, Gurney AL, Klijavin IU, Napier MA, Pan J, Paoni NF;  
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams EM, Wood WI;  
PI Zhang Z;  
XX  
XX WPI; 2003-102117/09.  
DR N-PSDB; ABX64020.  
XX  
PT Novel secreted and transmembrane polypeptide for modulating biological  
PT activity of cell expressing the polypeptide, identifying agonists or  
PT antagonists of polypeptide, and as molecular weight markers.  
XX  
PS Claim 12; Fig 68; 649pp; English.  
XX  
CC The present invention relates to the isolation of novel human PRO  
CC polypeptides, and the polynucleotide sequences encoding them. The PRO  
CC polypeptides are secreted and transmembrane proteins. The PRO  
CC polypeptides are useful for detecting other PRO polypeptides, for linking  
CC bioactive molecules to cells expressing PRO polypeptides, for modulating  
CC biological activities of cells expressing PRO polypeptides, and for for  
CC identifying agonists or antagonists. The polynucleotide sequences  
CC encoding PRO polypeptides are useful as hybridisation probes, in  
CC chromosome and gene mapping, in the generation of antisense RNA and DNA,  
CC in the preparation of PRO polypeptides, for generating transgenic animals  
CC or knockout animals, to construct hybridisation probes for mapping the  
CC gene which encodes the PRO polypeptide, and for the genetic analysis of  
CC individuals with genetic disorders, in gene therapy, for chromosome  
CC identification, as chromosome markers, and for generating probes for PCR,  
CC Northern analysis, Southern analysis and Western analysis. ABU13860-  
CC ABU14006 represent the human PRO polypeptides of the invention. Note: The  
CC sequence data for this patent was obtained in electronic format directly  
CC from the USPTO web site at seqdata.uspto.gov/psipsDIDEntry.html  
XX  
SQ Sequence 123 AA;  
Query Match 100.0%; Score 538; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 LDALLVFPQVQAQSCLTSPQHTVIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRADIP 60  
Dy 25 LDALLVFPQVQAQSCLTSPQHTVIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRADIP 84  
QY 61 DRFAAKDEAHNACVLTISPQEDDADYYCSVGFGFSP 99

DB 85 DRFAAKDEAHNACVLTISPQEDDADYYCSVGFGFSP 123  
RESULT 13  
ABU1070  
ID ABU1070 standard; protein; 123 AA.  
XX  
AC ABU1070;  
XX  
DT 23-JUN-2003 (first entry)  
XX  
DE Human PRO polypeptide #201.  
XX  
KW Human; PRO polypeptide; secreted and transmembrane protein;  
KW anti-PRO antibody; diagnostic assay; gene expression; diabetes;  
KW bone disorder; cartilage disorder; rheumatoid arthritis; obesity;  
KW sports injury; osteoarthritis; hyper-insulinaemia; hypo-insulinaemia;  
KW hearing loss; coagulation disorder; stroke; heart attack; cardiac;  
KW antidiabetic; anorectic; vulnery; antiarthritic; osteopathic;  
KW antirheumatic; auditory; cerebroprotective; angiogenic.  
XX  
OS Homo sapiens.  
XX  
FN US2003004311-A1.  
XX  
PD 02-JAN-2003.  
XX  
PF 19-DEC-2001; 2001US-00028072.  
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PR 18-JUN-1997; 97US-0049911P.  
PR 26-AUG-1997; 97US-0056974P.  
PR 17-SEP-1997; 97US-0059113P.  
PR 17-SEP-1997; 97US-0059115P.  
PR 17-SEP-1997; 97US-0059117P.  
PR 17-SEP-1997; 97US-0059122P.  
PR 17-SEP-1997; 97US-0059184P.  
PR 18-SEP-1997; 97US-0059263P.  
PR 18-SEP-1997; 97US-0059352P.  
PR 19-SEP-1997; 97US-0059588P.  
PR 24-SEP-1997; 97US-0059836P.  
PR 17-OCT-1997; 97US-0062250P.  
PR 17-OCT-1997; 97US-0062285P.  
PR 17-OCT-1997; 97US-0062287P.  
PR 24-OCT-1997; 97US-0063755P.  
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PR 24-OCT-1997; 97US-0062816P.  
PR 24-OCT-1997; 97US-0063045P.  
PR 24-OCT-1997; 97US-0063082P.  
PR 24-OCT-1997; 97US-0063127P.  
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PR 27-OCT-1997; 97US-0063329P.  
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PR 28-OCT-1997; 97US-0063561P.  
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PR 04-FEB-1998; 98US-0073612P.  
PR 09-FEB-1998; 98US-0074086P.

09-FEB-1998; 98US-0074032P.  
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27-MAR-1998; 98US-0079663P.  
27-MAR-1998; 98US-0079728P.  
31-MAR-1998; 98US-0080165P.  
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28-AUG-1998; 98US-0017888.  
10-SEP-1998; 98US-0018824.  
14-SEP-1998; 98US-0019093.  
14-SEP-1998; 98US-0019094.  
14-SEP-1998; 98US-0019177.  
16-SEP-1998; 98US-0019330.  
17-SEP-1998; 98US-0019437.  
07-OCT-1998; 98US-0021141.  
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29-OCT-1998; 98US-0022392.  
20-NOV-1998; 98US-0024855.  
01-DEC-1998; 98US-0025108.  
05-JAN-1999; 98US-0000106.  
08-MAR-1999; 98US-0005028.  
10-MAR-1999; 98US-0005190.  
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14-MAY-1999; 98US-0010733.  
02-JUN-1999; 98US-0012252.  
01-SEP-1999; 98US-0020111.  
08-SEP-1999; 98US-0020594.  
13-SEP-1999; 98US-0020944.  
15-SEP-1999; 98US-0021090.  
15-SEP-1999; 98US-0021347.  
05-OCT-1999; 98US-0023089.  
29-NOV-1999; 98US-0028214.  
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30-NOV-1999; 98US-0028409.  
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01-DEC-1999; 98US-0028634.  
02-DEC-1999; 98US-0028551.  
02-DEC-1999; 98US-0028564.  
02-DEC-1999; 98US-0028565.  
16-DEC-1999; 98US-0030095.  
20-DEC-1999; 98US-0030911.  
20-DEC-1999; 98US-0030999.  
30-DEC-1999; 98US-0031243.  
30-DEC-1999; 98US-0031274.  
05-JAN-2000; 2000US-0000219.  
06-JAN-2000; 2000US-0000277.  
11-FEB-2000; 2000US-000376.  
18-FEB-2000; 2000US-0003565.  
18-FEB-2000; 2000US-0004341.  
22-FEB-2000; 2000US-0004342.  
22-FEB-2000; 2000US-0004414.  
24-FEB-2000; 2000US-0004914.  
24-FEB-2000; 2000US-0005004.  
01-MAR-2000; 2000US-0005004.  
02-MAR-2000; 2000US-0005601.  
02-MAR-2000; 2000US-0005746.  
XX (GETH ) GENENTECH INC.  
XX Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;  
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;  
PI Smith V, Stewart TA, Tamas D, Watanabe CK, Wood WI, Zhang Z;  
DR WPI; 2003-352836/33.  
DR N-PSDB; ACA67194.  
XX  
XX New isolated PRO polypeptide useful for treating diabetes, rheumatoid  
PT arthritis, sports injuries, obesity, hearing loss in mammals, stroke, or  
PT heart attack.  
XX  
XX Claim 12; Fig 402; 643pp; English.

CC The present invention relates to the isolation of novel human PRO  
CC polypeptides, and the polynucleotide sequences encoding them. The PRO  
CC polypeptides are secreted and transmembrane proteins. The PRO  
CC polypeptides and polynucleotides are useful for preparing a medicament  
CC useful in the treatment of diabetes, bone and/or cartilage disorders  
CC (e.g. rheumatoid arthritis, sports injuries, osteoarthritis), obesity,  
CC hyper- or hypo-insulinaemia, hearing loss, and coagulation disorders  
CC (e.g. stroke, heart attack). Anti-PRO antibodies are useful in diagnostic  
CC assays for PRO, by detecting its expression in specific cells, tissues or  
CC serum, and for affinity purification of PRO from recombinant cell culture  
CC or natural sources. ABU0870-ABU81144 represent the human PRO  
CC polypeptides of the invention. Note: The sequence data for this patent  
CC was obtained in electronic format directly from the USPIO web site at  
CC seqdata.uspto.gov/psipdsIDentry.html  
XX  
XX Sequence 123 AA;  
SQ

Query Match 100.0%; Score 538; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51; Mismatches 0; Gaps 0;  
Matches 99; Conservative 0; Indels 0; Gaps 0;

QY 1 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIP 60  
Db 25 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIP 84  
QY 61 DRFSAKDEAHNACVLTISPQVEDDADYCVSGYGFSP 99  
Db 85 DRFSAKDEAHNACVLTISPQVEDDADYCVSGYGFSP 123

RESULT 14  
ABU72469  
ID ABU72469 standard; protein; 123 AA.  
XX  
XX ABU72469;  
XX  
XX 17-JUN-2003 (first entry)  
XX  
XX Novel human secreted and transmembrane protein PRO619.  
XX Human; secreted and transmembrane protein; cytostatic; anti-HIV;  
XX virucide; hepatotropic; antiinflammatory; neuroprotective; gene therapy;  
XX PRO; pharmaceutical; diagnostic; biosensor; bioindicator; malignancy;  
XX cancer; ovarian cancer; colorectal cancer; Kaposi's sarcoma; leukaemia;  
XX lymphoma; hepatitis B; multiple sclerosis; Crohn's disease;  
XX drug screening.  
XX Homo sapiens.  
XX  
XX US2003003531-A1.  
XX  
XX 02-JAN-2003.  
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XX 19-NOV-2001; 2001US-00989734.  
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XX 16-JUN-1997; 97US-0049787P.  
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XX 05-NOV-1997; 97US-0062250P.  
XX 12-NOV-1997; 97US-0065186P.  
XX 13-NOV-1997; 97US-0065311P.  
XX 24-NOV-1997; 97US-0066770P.  
XX 25-FEB-1998; 98US-0075945P.  
XX 20-MAR-1998; 98US-0078910P.  
XX 28-APR-1998; 98US-0083322P.  
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XX 28-MAY-1998; 98US-0087106P.  
XX 02-JUN-1998; 98US-0087607P.  
XX 02-JUN-1998; 98US-0087609P.  
XX 03-JUN-1998; 98US-0087759P.  
XX 03-JUN-1998; 98US-0087827P.  
XX 04-JUN-1998; 98US-0088021P.  
XX 04-JUN-1998; 98US-0088025P.  
XX 04-JUN-1998; 98US-0088026P.

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PR 04-JUN-1998; 98US-0088028P.
PR 04-JUN-1998; 98US-0088029P.
PR 04-JUN-1998; 98US-0088030P.
PR 04-JUN-1998; 98US-0088031P.
PR 04-JUN-1998; 98US-0088032P.
PR 05-JUN-1998; 98US-0088167P.
PR 05-JUN-1998; 98US-0088202P.
PR 05-JUN-1998; 98US-0088212P.
PR 05-JUN-1998; 98US-0088217P.
PR 05-JUN-1998; 98US-0088655P.
PR 09-JUN-1998; 98US-0088734P.
PR 10-JUN-1998; 98US-0088738P.
PR 10-JUN-1998; 98US-0088742P.
PR 10-JUN-1998; 98US-0088810P.
PR 10-JUN-1998; 98US-0088824P.
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PR 11-JUN-1998; 98US-0088858P.
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PR 11-JUN-1998; 98US-0088876P.
PR 12-JUN-1998; 98US-0089105P.
PR 16-JUN-1998; 98US-0089410P.
PR 16-JUN-1998; 98US-0089512P.
PR 16-JUN-1998; 98US-0089514P.
PR 17-JUN-1998; 98US-0089532P.
PR 17-JUN-1998; 98US-0089538P.
PR 17-JUN-1998; 98US-0089598P.
PR 17-JUN-1998; 98US-0089599P.
PR 17-JUN-1998; 98US-0089600P.
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PR 18-JUN-1998; 98US-0089801P.
PR 18-JUN-1998; 98US-0089907P.
PR 18-JUN-1998; 98US-0089917P.
PR 18-JUN-1998; 98US-0089908P.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 02-JUN-1999; 99WO-US012252.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
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PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 15-MAY-2000; 2000WO-US013358.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 28-JUL-2000; 2000WO-US015264.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 01-DEC-2000; 2000WO-US032678.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-JUN-2001; 2001WO-US017800.
PR 20-JUN-2001; 2001WO-US019692.
PR 29-JUN-2001; 2001WO-US021066.

PR 09-JUL-2001; 2001WO-US021735.
PR 28-AUG-2001; 2001US-00941992.
XX (GETH ) GENENTECH INC.
PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski RJ;
PI Grimaldi JC, Gurney AL, Kijavini IJ, Napier MA, Pan J, Paoni NF;
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;
PI Zhang Z;
XX WPI: 2003-352829/33.
DR N-PSDB; ACA64242.
XX New genes and secreted and transmembrane polypeptides (e.g. PRO183 or
PT PRO184), useful for treating or diagnosing e.g. ovarian cancer, Kaposi's
PT sarcoma, leukemia, lymphoma, hepatitis B, multiple sclerosis or Crohn's
PT disease.
XX Claim 12; Fig 68; 663pp; English.
XX The invention describes a new isolated nucleic acid molecule comprising
CC the full length coding sequence of the DNA deposited with the American
CC Type Culture Collection (e.g. ATCC Deposit No. 209621, 552-PTA, 819-PTA,
CC 209439, 203135, etc); or a sequence with at least 80% identity to a DNA
CC encoding a PRO polypeptide. The PRO polypeptides or polynucleotides are
CC useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These
CC are particularly useful for detecting or treating e.g. malignancies or
CC cancers (e.g. ovarian cancer, colorectal cancer, Kaposi's sarcoma,
CC leukemia or lymphoma). The PRO polypeptides are useful in drug screening,
CC disease in mammals. The PRO polypeptides are useful in screening,
CC particularly as targets for therapeutic intervention in these diseases,
CC and in the diagnostic determination of the presence of these diseases.
CC The PRO polypeptides are also useful as molecular weight markers, or for
CC chromosome identification. The PRO genes are useful as hybridisation
CC probes, or for screening libraries of human cDNA, genomic DNA or mRNA.
CC The PRO genes may also be used in gene therapy, particularly for
CC replacing a defective gene. This is the amino acid sequence of a novel
CC human secreted and transmembrane PRO polypeptide
XX Sequence 123 AA;
SQ Query Match 100.0%; Score 538; DB 6; Length 123;
Best Local Similarity 100.0%; Pred. No. 1e-51;
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIP 60
DB 25 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIP 84
QY 61 DRFSAKDEAHNACVLTITSPVQPEDDADYCSVGYGFSFSP 99
DB 85 DRFSAKDEAHNACVLTITSPVQPEDDADYCSVGYGFSFSP 123
RESULT 15
ABU66770
ID ABU66770 standard; protein; 123 AA.
XX AC ABU66770;
XX 23-MAY-2003 (first entry)
XX Human PRO polypeptide #201.
XX Human; PRO polypeptide; secreted and transmembrane protein;
KW tumour necrosis factor-alpha; TNF-alpha; blood; proliferation;
KW differentiation; chondrocyte; tumour; genetic disorder; cytostatic.
XX Homo sapiens.
XX US2003036180-A1.
XX
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PD 20-FEB-2003.  
XX 09-MAY-2002; 2002US-00143114.  
XX 31-MAR-1997; 97WO-US005230.  
PR 12-JUN-1998; 98WO-US012456.  
PR 14-JUL-1998; 98WO-US014552.  
PR 28-AUG-1998; 98WO-US017888.  
PR 10-SEP-1998; 98WO-US018824.  
PR 14-SEP-1998; 98WO-US019093.  
PR 14-SEP-1998; 98WO-US019177.  
PR 16-SEP-1998; 98WO-US019330.  
PR 17-SEP-1998; 98WO-US019437.  
PR 07-OCT-1998; 98WO-US021141.  
PR 29-OCT-1998; 98WO-US022991.  
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PR 05-JAN-1999; 98WO-US000106.  
PR 08-MAR-1999; 98WO-US0005028.  
PR 10-MAR-1999; 98WO-US0005190.  
PR 14-MAY-1999; 98WO-US0008615.  
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PR 10-MAR-2000; 2000WO-US006319.  
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PR 01-DEC-2000; 2000WO-US032678.  
PR 20-DEC-2000; 2000US-00747259.  
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PR 01-MAR-2001; 2001WO-US006666.  
PR 09-MAR-2001; 2001US-00802706.  
PR 14-MAR-2001; 2001US-00808689.  
PR 22-MAR-2001; 2001US-00816744.  
PR 05-APR-2001; 2001US-00828366.  
PR 10-MAY-2001; 2001US-00854208.  
PR 10-MAY-2001; 2001US-00854280.  
PR 18-MAY-2001; 2001US-00860216.  
PR 25-MAY-2001; 2001US-00866028.  
PR 25-MAY-2001; 2001US-00866034.  
PR 25-MAY-2001; 2001WO-US017092.  
PR 01-JUN-2001; 2001US-00872035.  
PR 01-JUN-2001; 2001WO-US017800.  
PR 05-JUN-2001; 2001US-00874503.  
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PR 19-JUN-2001; 2001US-00886342.  
PR 20-JUN-2001; 2001WO-US019692.  
PR 21-JUN-2001; 2001US-00887879.  
PR 22-JUN-2001; 2001WO-US020116.  
PR 29-JUN-2001; 2001WO-US021066.  
PR 09-JUL-2001; 2001WO-US021735.  
PR 18-JUL-2001; 2001US-00908827.  
PR 06-AUG-2001; 2001US-00924419.  
PR 09-AUG-2001; 2001US-00927796.  
PR 16-AUG-2001; 2001US-00931836.  
PR 19-DEC-2001; 2001US-00025072.  
XX (GETH ) GENENTECH INC.  
XX Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;  
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;  
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;  
XX WPI; 2003-332040/31.  
DR N-PSDB; ACA03803.  
XX New secreted and transmembrane PRO nucleic acids, useful for gene  
PT therapy, in chromosome and gene mapping, as chromosome markers, in tissue  
PT typing, and in chromosome identification.  
XX Claim 12; Fig 402; 660pp; English.  
XX The present invention relates to the isolation of novel human PRO  
CC polypeptides, and the polynucleotide sequences encoding them. The PRO  
CC polypeptides are secreted and transmembrane proteins. The PRO  
CC polypeptides are useful for detecting other PRO polypeptides, for linking  
CC bioactive molecules to cells expressing PRO polypeptides, for modulating  
CC biological activities of cells expressing PRO polypeptides, and for for  
CC identifying agonists or antagonists. The PRO polypeptides are useful for  
CC for stimulating the release of tumour necrosis factor (TNF)-alpha from  
CC human blood, for stimulating the proliferation or differentiation of  
CC chondrocytes, and detecting the presence of tumours. The polynucleotide  
CC sequences encoding PRO polypeptides are useful as hybridisation probes,  
CC in chromosome and gene mapping, in the generation of antisense RNA and  
CC DNA, in the preparation of PRO polypeptides, for generating transgenic  
CC animals or knockout animals, for the genetic analysis of individuals with  
CC genetic disorders, and in gene therapy. ABJ66570-ABJ66844 represent the  
CC human PRO polypeptides of the invention. Note: The sequence data for this  
CC patent was obtained in electronic format directly from the USPTO web site  
CC at seqdata.uspto.gov/psipsDIDEntry.html  
XX Sequence 123 AA;  
SQ  
Query Match 100.0%; Score 538; DB 6; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1e-51;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
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Db 25 LDALLVFEQVAQLSCTLSPQHVTIRDYGVSWYQQRAGSAPRYLLYYRSEEDHHRPADIP 84  
QY 61 DRFSAKDEAHNACVLTISPQPEDDADYVCVGVGFSP 99  
Db 85 DRFSAKDEAHNACVLTISPQPEDDADYVCVGVGFSP 123

Search completed: June 28, 2004, 08:26:33  
Job time : 46.4865 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: June 28, 2004, 08:28:50 ; Search time 35.2297 Seconds  
(without alignments)  
793.337 Million cell updates/sec

Title: US-09-981-876-200\_COPY\_25\_123

Perfect score: 538  
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Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1163542 seqs, 282313646 residues

Total number of hits satisfying chosen parameters: 1163542

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications AA:\*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
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2	538	100.0	123	9	US-09-989-723-117
3	538	100.0	123	9	US-09-989-727-117
4	538	100.0	123	9	US-09-989-727-117
5	538	100.0	123	9	US-09-989-731-117
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7	538	100.0	123	9	US-09-991-073-117
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9	538	100.0	123	9	US-09-991-163-117
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12	538	100.0	123	9	US-09-989-721-117
13	538	100.0	123	9	US-09-992-598-117
14	538	100.0	123	9	US-09-981-876-200
15	538	100.0	123	9	US-09-989-293A-117

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17	538	100.0	123	9	US-09-990-444-117
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19	538	100.0	123	9	US-09-989-730-117
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31	538	100.0	123	10	US-09-989-726-117
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35	538	100.0	123	10	US-09-997-514-117
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37	538	100.0	123	10	US-09-991-172-117
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39	538	100.0	123	10	US-09-997-559-117
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44	538	100.0	123	10	US-09-997-683-117
45	538	100.0	123	10	US-09-989-729A-117

ALIGNMENTS

RESULT 1

US-09-989-722-117

; Sequence 117, Application US/09989722

; Patent No. US20020072067A1

; GENERAL INFORMATION:

; APPLICANT: Ashkenazi, Avi J.

; APPLICANT: Baker, Kevin P.

; APPLICANT: Botstein, David

; APPLICANT: Desnoyers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Fong, Sherman

; APPLICANT: Gerber, Hanspeter

; APPLICANT: Gerritsen, Mary E.

; APPLICANT: Goddard, Audrey

; APPLICANT: Godowski, Paul J.

; APPLICANT: Grimaldi, J. Christopher

; APPLICANT: Gurney, Austin L.

; APPLICANT: Kljavin, Ivar J.

; APPLICANT: Napier, Mary A.

; APPLICANT: Pan, James

; APPLICANT: Paoni, Nicholas F.

; APPLICANT: Roy, Margaret Ann

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tamas, Daniel

; APPLICANT: Watanabe, Colin K.

; APPLICANT: Williams, P. Mickey

; APPLICANT: Wood, William I.

; APPLICANT: Zhang, Zemin

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

; TITLE OF INVENTION: Acids Encoding the Same

; FILE REFERENCE: P2730PIC63

; CURRENT APPLICATION NUMBER: US/09/989,722

; CURRENT FILING DATE: 2001-11-19

; PRIOR APPLICATION NUMBER: 60/049787

; PRIOR FILING DATE: 1997-06-16

; PRIOR APPLICATION NUMBER: 60/062250

; PRIOR FILING DATE: 1997-10-17

[illegible]

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; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091478
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091544
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match      100.0%; Score 538; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 1.9e-50;
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 25 LDALLVPGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYYRSEEDHHRPADIP 84
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Qy 61 DRFSAKDEAHNACVLTISVPQEDDADYGVSGYGFSP 99
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Db 85 DRFSAKDEAHNACVLTISVPQEDDADYGVSGYGFSP 123
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RESULT 2
US-09-989-723-117
; Sequence 117, Application US/09989723
; Patent No. US20020072092A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C62
; CURRENT APPLICATION NUMBER: US/09/989,723
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
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 ; PRIOR APPLICATION NUMBER: 60/091478

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 ; PRIOR APPLICATION NUMBER: 60/091544  
 ; PRIOR FILING DATE: 1998-07-01  
 ; PRIOR APPLICATION NUMBER: 60/091519  
 ; PRIOR FILING DATE: 1998-07-02  
 ; PRIOR APPLICATION NUMBER: 60/091626  
 ; PRIOR FILING DATE: 1998-07-02  
 ; PRIOR APPLICATION NUMBER: 60/091633  
 ; PRIOR FILING DATE: 1998-07-02  
 ; PRIOR APPLICATION NUMBER: 60/091978  
 ; PRIOR FILING DATE: 1998-07-07  
 ; PRIOR APPLICATION NUMBER: 60/091982  
 ; PRIOR FILING DATE: 1998-07-07  
 ; PRIOR APPLICATION NUMBER: 60/092182  
 ; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 538; DB 9; Length 123;  
 Best Local Similarity 100.0%; Pred. No. 1.9e-50;  
 Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LDALLVFGQVAQLSCTLSPOHVTIRYGVSWYQQRAGSAPRYLLYYRSEEDHHRPADIP 60  
 Db 25 LDALLVFGQVAQLSCTLSPOHVTIRYGVSWYQQRAGSAPRYLLYYRSEEDHHRPADIP 84  
 QY 61 DRFSAKDEAHNACVLTISPVQEDDADYCSGVGYGFSF 99  
 Db 85 DRFSAKDEAHNACVLTISPVQEDDADYCSGVGYGFSF 123

RESULT 3  
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 ; Sequence 117, Application US/09989279  
 ; Patent No. US20020072496A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Ashkenazi, Avi J.  
 ; APPLICANT: Baker, Kevin P.  
 ; APPLICANT: Botstein, David  
 ; APPLICANT: Desnoyers, Luc  
 ; APPLICANT: Eaton, Dan L.  
 ; APPLICANT: Ferrara, Napoleone  
 ; APPLICANT: Fong, Sherman  
 ; APPLICANT: Gerber, Hanspeter  
 ; APPLICANT: Gerritsen, Mary E.  
 ; APPLICANT: Goddard, Audrey  
 ; APPLICANT: Godowski, Paul J.  
 ; APPLICANT: Grimaldi, J. Christopher  
 ; APPLICANT: Gurney, Austin L.  
 ; APPLICANT: Kljavin, Ivar J.  
 ; APPLICANT: Napier, Mary A.  
 ; APPLICANT: Pan, James  
 ; APPLICANT: Paoni, Nicholas F.  
 ; APPLICANT: Roy, Margaret Ann  
 ; APPLICANT: Stewart, Timothy A.  
 ; APPLICANT: Tumas, Daniel  
 ; APPLICANT: Watanabe, Colin K.  
 ; APPLICANT: Williams, P. Mickey  
 ; APPLICANT: Wood, William I.  
 ; APPLICANT: Zhang, Zemin  
 ; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 ; TITLE OF INVENTION: Acids Encoding the Same  
 ; FILE REFERENCE: P2730PIC56  
 ; CURRENT APPLICATION NUMBER: US/09/989,279  
 ; CURRENT FILING DATE: 2001-11-19  
 ; PRIOR APPLICATION NUMBER: 60/049787  
 ; PRIOR FILING DATE: 1997-06-16  
 ; PRIOR APPLICATION NUMBER: 60/062250  
 ; PRIOR FILING DATE: 1997-10-17  
 ; PRIOR APPLICATION NUMBER: 60/065186  
 ; PRIOR FILING DATE: 1997-11-12  
 ; PRIOR APPLICATION NUMBER: 60/065311  
 ; PRIOR FILING DATE: 1997-11-13  
 ; PRIOR APPLICATION NUMBER: 60/065770  
 ; PRIOR FILING DATE: 1997-11-24

[illegible]

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; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match      100.0%; Score 538; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 1.9e-50;
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEEDHHRPADIP 60
   |||||
Db 25 LDALLVFGQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEEDHHRPADIP 84
   |||||

QY 61 DRFSAKDDEAHNACVLITSPVQPEDDADYCVSGVGFSP 99
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Db 85 DRFSAKDDEAHNACVLITSPVQPEDDADYCVSGVGFSP 123

RESULT 4
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; Sequence 117, Application US/09989727
; Patent No. US20020072497A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Nepier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C65
; CURRENT APPLICATION NUMBER: US/09/989, 727
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
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; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match      100.0%; Score 538; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 1.9e-50;
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 2S LDALLVPPGQVAQLSCTLSFQHVITRDYGVSWTQQRAGSAPRYLLYYRSEEDHHRPADIP 84

Qy 61 DRFSAKDEAHNACVLITISVPQEDDADYICSVGYGFSP 99
Db 85 DRFSAKDEAHNACVLITISVPQEDDADYICSVGYGFSP 123

RESULT 5
US-09-989-731-117
; Sequence 117, Application US/09989731
; Patent No. US20020103125A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kijavini, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C70
; CURRENT APPLICATION NUMBER: US/09/989,731
; CURRENT FILING DATE: 2001-11-20
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; PRIOR FILING DATE: 1998-04-28

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; PRIOR FILING DATE: 1998-07-09

Query Match      100.0%; Score 538; DB 9; Length 123;
Best Local Similarity 100.0%; Pred. No. 1.9e-50;
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 25 LDALLVPPGVAQLSCTLSQHTVIRDYGVSWYQQAGAPRYLLLYRSEEDHHRPADIP 84

Qy 61 DRFSAKDEAHNAACVLITISVPQEDDADYVCVGYGFSP 99
Db 85 DRFSAKDEAHNAACVLITISVPQEDDADYVCVGYGFSP 123

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; Patent No. US20020123463A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrari, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
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; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C57
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; PRIOR APPLICATION NUMBER: 60/090676  
; PRIOR FILING DATE: 1998-06-25  
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; PRIOR FILING DATE: 1998-06-25  
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; PRIOR FILING DATE: 1998-07-02  
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; PRIOR APPLICATION NUMBER: 60/091633  
; PRIOR FILING DATE: 1998-07-02  
; PRIOR APPLICATION NUMBER: 60/091978  
; PRIOR FILING DATE: 1998-07-07  
; PRIOR APPLICATION NUMBER: 60/091982

; PRIOR FILING DATE: 1998-07-07  
; PRIOR APPLICATION NUMBER: 60/092182  
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 538; DB 9; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1.9e-50;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LDALLVFPQVAQLSCTLSPOHVTIRDVGSWYQCRAGSAPRYLLYYRSEDDHHRPADIP 60

DB 25 LDALLVFPQVAQLSCTLSPOHVTIRDVGSWYQCRAGSAPRYLLYYRSEDDHHRPADIP 84

QY 61 DRFSAKDEAHNACVLTISVPQPEDDADYYCSVGYGFSF 99

DB 85 DRFSAKDEAHNACVLTISVPQPEDDADYYCSVGYGFSF 123

## RESULT 7

US-09-991-073-117  
; Sequence 117, Application US/09991073  
; Patent No. US20020127576A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnoyers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gersitsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, J. Christopher  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
; APPLICANT: Pan, James  
; APPLICANT: Paoni, Nicholas F.  
; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Williams, P. Mickey  
; APPLICANT: Wood, William I.  
; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
; TITLE OF INVENTION: Acids Encoding the Same  
; FILE REFERENCE: P2730PIC15  
; CURRENT APPLICATION NUMBER: US/09/991,073  
; PRIOR FILING DATE: 2001-11-14  
; PRIOR APPLICATION NUMBER: 60/049787  
; PRIOR FILING DATE: 1997-06-16  
; PRIOR APPLICATION NUMBER: 60/062250  
; PRIOR FILING DATE: 1997-10-17  
; PRIOR APPLICATION NUMBER: 60/065186  
; PRIOR FILING DATE: 1997-11-12  
; PRIOR APPLICATION NUMBER: 60/065311  
; PRIOR FILING DATE: 1997-11-13  
; PRIOR APPLICATION NUMBER: 60/066770  
; PRIOR FILING DATE: 1997-11-24  
; PRIOR APPLICATION NUMBER: 60/075945  
; PRIOR FILING DATE: 1998-02-25  
; PRIOR APPLICATION NUMBER: 60/078910  
; PRIOR FILING DATE: 1998-03-20  
; PRIOR APPLICATION NUMBER: 60/083322  
; PRIOR FILING DATE: 1998-04-28  
; PRIOR APPLICATION NUMBER: 60/084600  
; PRIOR FILING DATE: 1998-05-07  
; PRIOR APPLICATION NUMBER: 60/087106  
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; PRIOR FILING DATE: 1998-06-02

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Query Match 100.0%; Score 538; DB 9; Length 123;  
Best Local Similarity 100.0%; Pred.No. 1.9e-50;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LDALLVPPGVAQLSCTLSQHTVIRYGVSWYQQRAGSAPRYLLYYRSEEDHHRPADIP 60  
Db 25 LDALLVPPGVAQLSCTLSQHTVIRYGVSWYQQRAGSAPRYLLYYRSEEDHHRPADIP 84

Qy 61 DRSAAKDEAHNAVLITISVQEDDADYKCSVGYGSP 99  
Db 85 DRSAAKDEAHNAVLITISVQEDDADYKCSVGYGSP 123

RESULT 8  
US-09-990-442-117  
; Sequence 117, Application US/09990442  
; Patent No. US20020132252A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Deanoyers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, J. Christopher  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
; APPLICANT: Pan, James  
; APPLICANT: Paoni, Nicholas F.  
; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Williams, P. Mickey  
; APPLICANT: Wood, William I.  
; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
; TITLE OF INVENTION: Acids Encoding the Same  
; FILE REFERENCE: P2730PIC8  
; CURRENT APPLICATION NUMBER: US/09/990,442  
; CURRENT FILING DATE: 2001-11-14  
; PRIOR APPLICATION NUMBER: 60/049787  
; PRIOR FILING DATE: 1997-06-16  
; PRIOR APPLICATION NUMBER: 60/062250  
; PRIOR FILING DATE: 1997-10-17  
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; PRIOR FILING DATE: 1997-11-24  
; PRIOR APPLICATION NUMBER: 60/075945  
; PRIOR FILING DATE: 1998-02-25  
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; PRIOR FILING DATE: 1998-03-20  
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; PRIOR FILING DATE: 1998-06-02  
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; PRIOR APPLICATION NUMBER: 60/087759

; PRIOR FILING DATE: 1998-06-02  
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; PRIOR FILING DATE: 1998-06-03  
; PRIOR APPLICATION NUMBER: 60/088021  
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; PRIOR FILING DATE: 1998-06-04  
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;; PRIOR FILING DATE: 1998-07-01  
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;; PRIOR FILING DATE: 1998-07-01  
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;; PRIOR FILING DATE: 1998-07-02  
;; PRIOR APPLICATION NUMBER: 60/091633  
;; PRIOR FILING DATE: 1998-07-02  
;; PRIOR APPLICATION NUMBER: 60/091978  
;; PRIOR FILING DATE: 1998-07-07  
;; PRIOR APPLICATION NUMBER: 60/091982  
;; PRIOR FILING DATE: 1998-07-07  
;; PRIOR APPLICATION NUMBER: 60/092182  
;; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 538; DB 9; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1.9e-50;

Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 LDALLVFCQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIP 60  
Db 25 LDALLVFCQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHRRPADIP 84  
Qy 61 DRFSAKDEAHNACVLTISPQVEDDADYCVSGVGFSP 99  
Db 85 DRFSAKDEAHNACVLTISPQVEDDADYCVSGVGFSP 123

RESULT 9

US-09-991-163-117  
; Sequence 117, Application US/09991163  
; Patent No. US20020132253A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnoyers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
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; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
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; APPLICANT: Pan, James  
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; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Williams, P. Mickey  
; APPLICANT: Wood, William I.  
; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
; FILE REFERENCE: P2730PIC17  
; CURRENT APPLICATION NUMBER: US/09/991,163  
; CURRENT FILING DATE: 2001-11-14  
; PRIOR APPLICATION NUMBER: 60/049787  
; PRIOR FILING DATE: 1997-06-16  
; PRIOR APPLICATION NUMBER: 60/062250  
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 ; PRIOR FILING DATE: 1998-06-04  
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; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnovers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferrara, Napoleone  
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; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
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; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Williams, P. Mickey  
; APPLICANT: Wood, William I.  
; APPLICANT: Zhang, Zemin  
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Sequence 117 Application US/09990456  
Patent No. US20020137890A1  
GENERAL INFORMATION:  
APPLICANT: Ashkenazi, Avi J.  
APPLICANT: Baker, Kevin P.  
APPLICANT: Botstein, David  
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APPLICANT: Zhang, Zemin  
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
FILE REFERENCE: P27301C22  
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; PRIOR APPLICATION NUMBER: 60/091982  
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; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 538; DB 9; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1.9e-50;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LDALLVPGVQVACLCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADLP 60  
Db 25 LDALLVPGVQVACLCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYRSEEDHHRPADLP 84  
Qy 61 DRFSAKDEAHNACVLTISFVQPEDDADYVCVGVGFSP 99  
Db 85 DRFSAKDEAHNACVLTISFVQPEDDADYVCVGVGFSP 123

## RESULT 12

US-09-989-721-117  
; Sequence 117, Application US/09989721  
; Patent No. US20020142961A1  
; GENERAL INFORMATION:  
; APPLICANT: Ashkenazi, Avi J.  
; APPLICANT: Baker, Kevin P.  
; APPLICANT: Botstein, David  
; APPLICANT: Desnovers, Luc  
; APPLICANT: Eaton, Dan L.  
; APPLICANT: Ferraro, Napoleone  
; APPLICANT: Fong, Sherman  
; APPLICANT: Gerber, Hanspeter  
; APPLICANT: Gerritsen, Mary E.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, J. Christopher  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Kljavin, Ivar J.  
; APPLICANT: Napier, Mary A.  
; APPLICANT: Pan, James  
; APPLICANT: Paoni, Nicholas F.  
; APPLICANT: Roy, Margaret Ann  
; APPLICANT: Stewart, Timothy A.  
; APPLICANT: Tumas, Daniel  
; APPLICANT: Watanabe, Colin K.  
; APPLICANT: Williams, P. Mickey  
; APPLICANT: Wood, William I.  
; APPLICANT: Zhang, Zemin  
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
; FILE REFERENCE: P2730P1C55  
; CURRENT APPLICATION NUMBER: US/09/989,721  
; CURRENT FILING DATE: 2001-11-19  
; PRIOR APPLICATION NUMBER: 60/049787  
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PRIOR FILING DATE: 1998-07-07  
PRIOR APPLICATION NUMBER: 60/091982  
PRIOR FILING DATE: 1998-07-07  
PRIOR APPLICATION NUMBER: 60/092182  
PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 538; DB 9; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1.9e-50;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LDALLVFGQVAQSLTSPDHVTRDYGVSWYQQRAGSAPRYLLYYRSEDDHRRPADIP 60  
Db 25 LDALLVFGQVAQSLTSPDHVTRDYGVSWYQQRAGSAPRYLLYYRSEDDHRRPADIP 84  
Qy 61 DRFSAKDEAHNACVLITSPQPDADYYCSVGYGFSF 99  
Db 85 DRFSAKDEAHNACVLITSPQPDADYYCSVGYGFSF 123

RESULT 13

US-09-992-598-117  
Sequence 117, Application US/09992598  
Patent No. US20020160384A1  
GENERAL INFORMATION:  
APPLICANT: Ashkenazi, Avi J.  
APPLICANT: Baker, Kevin P.  
APPLICANT: Botstein, David  
APPLICANT: Desnoyers, Luc  
APPLICANT: Eaton, Dan L.  
APPLICANT: Ferrara, Napoleone  
APPLICANT: Fong, Sherman  
APPLICANT: Gerber, Hanspeter  
APPLICANT: Gottfredsen, Mary E.  
APPLICANT: Goddard, Audrey  
APPLICANT: Godowski, Paul J.  
APPLICANT: Grimaldi, J. Christopher  
APPLICANT: Gurney, Austin L.  
APPLICANT: Kljavin, Ivar J.  
APPLICANT: Napier, Mary A.  
APPLICANT: Pan, James  
APPLICANT: Paoni, Nicholas F.  
APPLICANT: Roy, Margaret Ann  
APPLICANT: Stewart, Timothy A.  
APPLICANT: Tumas, Daniel  
APPLICANT: Watanabe, Colin K.  
APPLICANT: Williams, P. Mickey  
APPLICANT: Wood, William I.  
APPLICANT: Zhang, Zemin  
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
TITLE OF INVENTION: Acids Encoding the Same  
FILE REFERENCE: P2730PIC20  
CURRENT APPLICATION NUMBER: US/09/992,598  
CURRENT FILING DATE: 2001-11-14  
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;	PRIOR FILING DATE:	1998-07-07
;	PRIOR APPLICATION NUMBER:	60/091982
;	PRIOR FILING DATE:	1998-07-07
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;	PRIOR FILING DATE:	1998-07-09

Query Match 100.0%; Score 538; DB 9; Length 123;  
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Qy	1	LDALLPFCGVQLSCTTSFQHVTTIRDYGVSVYQQRAGSAPRYLLLYRSEEDHHPADIP	60
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Qy	61	DRFSAKDEAHNACVLTTISFVQPEDDADYCVSVGYGFSF	99
Db	85	DRFSAKDEAHNACVLTTISFVQPEDDADYCVSVGYGFSF	123

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; Sequence 200, Application US/09981876  
; Patent No. US20020164669A1

GENERAL INFORMATION:  
; APPLICANT: Rosen et al.  
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; FILE REFERENCE: P2001P1  
; CURRENT APPLICATION NUMBER: US/09/981,876  
; CURRENT FILING DATE: 2001-10-19  
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; PRIOR FILING DATE: 1997-04-11  
; PRIOR APPLICATION NUMBER: 60/043,315  
; PRIOR FILING DATE: 1997-04-11  
; PRIOR APPLICATION NUMBER: 60/048,974  
; PRIOR FILING DATE: 1997-06-06  
; PRIOR APPLICATION NUMBER: 60/056,886  
; PRIOR FILING DATE: 1997-08-22  
; PRIOR APPLICATION NUMBER: 60/056,877  
; PRIOR FILING DATE: 1997-08-22  
; PRIOR APPLICATION NUMBER: 60/056,889  
; PRIOR FILING DATE: 1997-08-22  
; PRIOR APPLICATION NUMBER: 60/056,893  
; PRIOR FILING DATE: 1997-08-22  
; PRIOR APPLICATION NUMBER: 60/056,630  
; PRIOR FILING DATE: 1997-08-22  
; PRIOR APPLICATION NUMBER: 60/056,878  
; PRIOR FILING DATE: 1997-08-22  
; PRIOR APPLICATION NUMBER: 60/056,662  
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; PRIOR FILING DATE: 1997-08-22  
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; PRIOR FILING DATE: 1997-08-22  
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; PRIOR APPLICATION NUMBER: 60/047,595  
; PRIOR FILING DATE: 1997-05-23  
; PRIOR APPLICATION NUMBER: 60/057,761  
; PRIOR FILING DATE: 05-SEP-1997  
; PRIOR APPLICATION NUMBER: 60/047,599  
; PRIOR FILING DATE: 1997-05-23

PRIOR APPLICATION NUMBER: 60/047,588  
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PRIOR FILING DATE: 1997-05-23  
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PRIOR APPLICATION NUMBER: 60/043,576  
PRIOR FILING DATE: 1997-04-11  
PRIOR APPLICATION NUMBER: 60/047,501  
PRIOR FILING DATE: 1997-05-23  
PRIOR APPLICATION NUMBER: 60/043,670  
PRIOR FILING DATE: 1997-04-11  
PRIOR APPLICATION NUMBER: 60/056,632  
PRIOR FILING DATE: 1997-08-22  
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PRIOR FILING DATE: 1997-08-22  
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PRIOR FILING DATE: 1997-08-22  
PRIOR APPLICATION NUMBER: 60/056,908  
PRIOR FILING DATE: 1997-08-22  
PRIOR APPLICATION NUMBER: 60/048,964  
PRIOR FILING DATE: 1997-06-06  
PRIOR APPLICATION NUMBER: 60/057,650  
PRIOR FILING DATE: 1997-09-05  
PRIOR APPLICATION NUMBER: 60/056,884  
PRIOR FILING DATE: 1997-08-22  
NUMBER OF SEQ ID NOS: 280  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 200  
LENGTH: 123

Query Match 100.0%; Score 538; DB 9; Length 123;  
Best Local Similarity 100.0%; Pred. No. 1.9e-50;  
Matches 99; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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DB 25 LDALLVFGQVAQLSCTLSPQHVIRDYGVSWYQQRAGSAPRYLLYYRSEEDHRRPADIP 84  
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DB 85 DRFSAKDDEAHNACVLITSPQVEDDADYCVSGYGFSP 123

RESULT 15

US-09-989-293A-117  
Sequence 117, Application US/09989293A  
Patent No. US2002017164A1  
GENERAL INFORMATION:  
APPLICANT: Ashkenazi, Avi J.  
APPLICANT: Baker, Kevin P.

APPLICANT: Botstein, David  
APPLICANT: Desnoyers, Luc  
APPLICANT: Eaton, Dan L.  
APPLICANT: Ferrara, Napoleone  
APPLICANT: Fong, Sherman  
APPLICANT: Gerber, Hanspeter  
APPLICANT: Gerritsen, Mary E.  
APPLICANT: Goddard, Audrey  
APPLICANT: Godowski, Paul J.  
APPLICANT: Grimaldi, J. Christopher  
APPLICANT: Gurney, Austin L.  
APPLICANT: Kljavin, Ivar J.  
APPLICANT: Napier, Mary A.  
APPLICANT: Pan, James  
APPLICANT: Paoni, Nicholas F.  
APPLICANT: Roy, Margaret Ann  
APPLICANT: Stewart, Timothy A.  
APPLICANT: Tumas, Daniel  
APPLICANT: Watanabe, Colin K.  
APPLICANT: Williams, P. Mickey  
APPLICANT: Wood, William I.  
APPLICANT: Zhang, Zemin  
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
FILE REFERENCE: P2730P1C66  
CURRENT APPLICATION NUMBER: US/09/989,293A  
CURRENT FILING DATE: 2001-11-20  
PRIOR APPLICATION NUMBER: 60/049787  
PRIOR FILING DATE: 1997-06-16  
PRIOR APPLICATION NUMBER: 60/062250  
PRIOR FILING DATE: 1997-10-17  
PRIOR APPLICATION NUMBER: 60/065186  
PRIOR FILING DATE: 1997-11-12  
PRIOR APPLICATION NUMBER: 60/065311  
PRIOR FILING DATE: 1997-11-13  
PRIOR APPLICATION NUMBER: 60/066770  
PRIOR FILING DATE: 1997-11-24  
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PRIOR FILING DATE: 1998-02-25  
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PRIOR FILING DATE: 1998-03-20  
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PRIOR FILING DATE: 1998-04-28  
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PRIOR FILING DATE: 1998-05-07  
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PRIOR FILING DATE: 1998-06-04  
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PRIOR FILING DATE: 1998-06-04  
PRIOR APPLICATION NUMBER: 60/088033  
PRIOR FILING DATE: 1998-06-04  
PRIOR APPLICATION NUMBER: 60/088326  
PRIOR FILING DATE: 1998-06-04  
PRIOR APPLICATION NUMBER: 60/088167  
PRIOR FILING DATE: 1998-06-05







GenCore version 5.1.6  
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OM protein - protein search, using sw model

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Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

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Post-processing: Minimum Match 0%  
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Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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4	189	35.1	110	2	US-08-362-780-16
5	188.5	35.0	249	4	US-08-918-148-74
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7	182.5	33.9	245	4	US-10-039-785-42
8	181.5	33.7	110	3	US-09-240-274-63
9	181	33.6	107	4	US-09-025-769B-34
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19	174.5	32.4	235	3	US-09-049-672A-10
20	174.5	32.4	244	4	US-08-918-148-79
21	173.5	32.2	106	1	US-08-488-113B-155
22	173.5	32.2	106	1	US-08-477-494B-155
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24	173.5	32.2	106	1	US-08-472-788A-19
25	173.5	32.2	106	2	US-08-477-531B-19
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27	173.5	32.2	106	2	US-08-082-842A-19

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37	173	32.2	112	4	US-09-315-574-39
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44	171	31.8	97	2	US-08-665-202-35
45	171	31.8	97	4	US-09-315-574-35

ALIGNMENTS

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; Sequence 200, Application US/09148545  
; Patent No. 6590075  
; GENERAL INFORMATION:  
; APPLICANT: Rosen et al.  
; TITLE OF INVENTION: 70 Human Secreted Proteins  
; FILE REFERENCE: PZ001PI  
; CURRENT APPLICATION NUMBER: US/09/148,545  
; CURRENT FILING DATE: 1998-09-04  
; EARLIER APPLICATION NUMBER: PCT/US98/04482  
; EARLIER FILING DATE: 1998-03-06  
; EARLIER APPLICATION NUMBER: 60/040,162  
; EARLIER FILING DATE: 1997-03-07  
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; EARLIER FILING DATE: 1997-03-07  
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; EARLIER APPLICATION NUMBER: 60/047,581  
; EARLIER FILING DATE: 1997-05-23



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; EARLIER FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 280
; SOFTWARE: Patent in Ver. 2.0
; SEQ ID NO 200
; LENGTH: 123

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Db 25 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEEDHRRPADIP 84

Qy 61 DRFSAKDEAHNACVLTISVPQEDDADYYCSVGYGFSF 99
Db 85 DRFSAKDEAHNACVLTISVPQEDDADYYCSVGYGFSF 123

RESULT 2
US-09-621-976-5367
; Sequence 5367, Application US/09621976
; Patent No. 6639063
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Jobert, S.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: ESTs and Encoded Human Proteins.
; FILE REFERENCE: GENSET.054PR2
; CURRENT APPLICATION NUMBER: US/09/621,976
; CURRENT FILING DATE: 2000-07-21
; NUMBER OF SEQ ID NOS: 19335
; SOFTWARE: Patent.pm
; SEQ ID NO 5367
; LENGTH: 123
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; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SIGNAL
; LOCATION: -20...-1
US-09-621-976-5367

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Db 25 LDALLVFPQVAQLSCTLSPOHVTIRDYGVSWYQORAGSAPRYLLYYRSEEDHRRPADIP 84

Qy 61 DRFSAKDEAHNACVLTISVPQEDDADYYCSVGYGFSF 99
Db 85 DRFSAKDEAHNACVLTISVPQEDDADYYCSVGYGFSF 123

RESULT 3
US-07-988-925-16
; Sequence 16, Application US/07988925
; Patent No. 5585097
; GENERAL INFORMATION:
; APPLICANT: Bolt, Sarah L.
; APPLICANT: Clark, Michael R.
; APPLICANT: Gorman, Scott D.
; APPLICANT: Routledge, Edward G.
; APPLICANT: Waldmann, Herman
; TITLE OF INVENTION: antibody preparation
; NUMBER OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Nixon and Vanderhye pc
; STREET: 11th Floor, 1100 No. 5585097th Glebe Road
; CITY: Arlington
; STATE: Virginia
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; COUNTRY: USA
; ZIP: 22201
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/988,925
; FILING DATE:
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: GB 9206422.9
; FILING DATE: 24-MAR-1992
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: WO PCT/GB92/01933
; FILING DATE: 21-OCT-1992
; ATTORNEY/AGENT INFORMATION:
; NAME: Mitchard, Leonard C
; REGISTRATION NUMBER: 29009
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 7038164000
; TELEFAX: 7038164100
; INFORMATION FOR SEQ ID NO: 16:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 110 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-07-988-925-16

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Matches 39; Conservative 14; Mismatches 25; Indels 6; Gaps 2;

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Db 14 PGKTVISCTLSGN--ENNYVHWYQRPGRPTTVF-----DDKRPDGVDFRPSGI 67

Qy 68 DEAHNACVLTISVPQEDDADYYC 91
Db 68 DRSSNSASLTISGLQTEDEADYYC 91

RESULT 4
US-08-362-780-16
; Sequence 16, Application US/08362780
; Patent No. 5968509
; GENERAL INFORMATION:
; APPLICANT: Gorman, Scott D.
; APPLICANT: Routledge, Edward G.
; APPLICANT: Waldmann, Herman
; TITLE OF INVENTION: Antibody Preparation
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Nixon and Vanderhye pc
; STREET: 9th Floor, 1100 No. 5968509th Glebe Road
; CITY: Arlington
; STATE: Virginia
; COUNTRY: USA
; ZIP: 22201
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/362,780
; FILING DATE:
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/862,543
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Matches 39; Conservative 15; Mismatches 24; Indels 7; Gaps 3;  
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Qy 68 DEAHNACVLITSPVQPEDADYYCS 92  
Db 203 --SGNTASLTSGVGLQAEADYYCS 225  
RESULT 8  
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; Sequence 63, Application US/09240274  
; Patent No. 6255455  
; GENERAL INFORMATION:  
; APPLICANT: Siegel, Donald L.  
; TITLE OF INVENTION: RH(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL  
; TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF  
; FILE REFERENCE: 09596-42U2  
; CURRENT APPLICATION NUMBER: US/09/240,274  
; CURRENT FILING DATE: 1999-01-29  
; EARLIER APPLICATION NUMBER: 60/081,380  
; EARLIER FILING DATE: 1998-04-10  
; EARLIER APPLICATION NUMBER: 60/028,550  
; EARLIER FILING DATE: 1996-10-11  
; NUMBER OF SEQ ID NOS: 224  
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; ORGANISM: Homo sapiens  
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US-09-240-274-63

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Qy 68 DEAHNACVLITSPVQPEDADYYC 91  
Db 68 --SGTSASLAITGQAEADYYC 89

RESULT 9  
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; Sequence 34, Application US/09025769B  
; Patent No. 6300064  
; GENERAL INFORMATION:  
; APPLICANT: Knappik, Achim  
; APPLICANT: Pack, Peter  
; APPLICANT: Ilag, Vic  
; APPLICANT: Ge, Liming  
; APPLICANT: Moroney, Simon  
; APPLICANT: Plueckthun, Andreas  
; TITLE OF INVENTION: Protein/(Poly)peptide libraries  
; NUMBER OF SEQUENCES: 373  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave  
; STREET: 1251 Avenue of the Americas  
; CITY: New York  
; STATE: New York  
; COUNTRY: USA  
; ZIP: 10021  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: IBM PC compatible  
; SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)  
; CURRENT APPLICATION NUMBER: US/09/025,769B  
; FILING DATE: 18-FEB-1998  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: EP 95 11 3021.0  
; FILING DATE: 18-AUG-1995  
; ATTORNEY/AGENT INFORMATION:  
; NAME: James F. Haley, Jr., Esq.  
; REGISTRATION NUMBER: 27,794  
; REFERENCE/DOCKET NUMBER: MORPHO/5  
; TELECOMMUNICATION INFORMATION:

; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/025,769B  
; FILING DATE: 18-FEB-1998  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: EP 95 11 3021.0  
; FILING DATE: 18-AUG-1995  
; ATTORNEY/AGENT INFORMATION:  
; NAME: James F. Haley, Jr., Esq.  
; REGISTRATION NUMBER: 27,794  
; REFERENCE/DOCKET NUMBER: MORPHO/5  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (212)596-9000  
; TELEFAX: (212)596-9090  
; INFORMATION FOR SEQ ID NO: 34:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 107 amino acids  
; TYPE: amino acid  
; STRANDEDNESS:  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; US-09-025-769B-34  
Query Match 33.6%; Score 181; DB 4; Length 107;  
Best Local Similarity 42.6%; Pred. No. 8.1e-13;  
Matches 40; Conservative 13; Mismatches 31; Indels 10; Gaps 3;  
Qy 6 VFGQVAQLSCTLSPOHVTIRYGVSWYQORAGSAPRYLLYRSEEDHHRPADIDRFSAA 65  
Db 12 VAPGQTARISCSGD---ALGDKYASWYQKPGQAPLVLY----DDSRPSGIDRFSG 63  
Qy 66 AKDEAHNACVLITSPVQPEDADYYCSVGYGFSP 99  
Db 64 S--NSCNTATLTISGTQAEADYYCQHHYTPP 95

RESULT 10  
US-09-025-769B-55  
; Sequence 55, Application US/09025769B  
; Patent No. 6300064  
; GENERAL INFORMATION:  
; APPLICANT: Knappik, Achim  
; APPLICANT: Pack, Peter  
; APPLICANT: Ilag, Vic  
; APPLICANT: Ge, Liming  
; APPLICANT: Moroney, Simon  
; APPLICANT: Plueckthun, Andreas  
; TITLE OF INVENTION: Protein/(Poly)peptide libraries  
; NUMBER OF SEQUENCES: 373  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave  
; STREET: 1251 Avenue of the Americas  
; CITY: New York  
; STATE: New York  
; COUNTRY: USA  
; ZIP: 10021  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: IBM PC compatible  
; SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)  
; CURRENT APPLICATION NUMBER: US/09/025,769B  
; FILING DATE: 18-FEB-1998  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: EP 95 11 3021.0  
; FILING DATE: 18-AUG-1995  
; ATTORNEY/AGENT INFORMATION:  
; NAME: James F. Haley, Jr., Esq.  
; REGISTRATION NUMBER: 27,794  
; REFERENCE/DOCKET NUMBER: MORPHO/5  
; TELECOMMUNICATION INFORMATION:

TELEPHONE: (212)596-9000  
 TELEFAX: (212)596-9090  
 INFORMATION FOR SEQ ID NO: 55:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 107 amino acids  
 TYPE: amino acid  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-09-025-769B-55

Query Match 33.6%; Score 181; DB 4; Length 107;  
 Best Local Similarity 42.6%; Pred. No. 8.1e-13;  
 Matches 40; Conservative 13; Mismatches 31; Indels 10; Gaps 3;  
 QY 6 VFGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSA 65  
 DB 12 VAPQOTARISCGD----ALGDKYASWYQXPGQAPVLVIY----DSDRPSGIPERFSG 63  
 QY 66 AKDEAHNACVLITISVPQEDDADYCVSGYGFSP 99  
 DB 64 S--NSGNTATLTISGTAQDEADYCCQHYTTPP 95

RESULT 11  
 US-10-039-785-48  
 ; Sequence 48, Application US/10039785  
 ; Patent No. 6538938  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Salcedo et al.  
 ; TITLE OF INVENTION: Antibodies that Immunospecifically Bind to TRAIL  
 ; FILE REFERENCE: PF550  
 ; CURRENT APPLICATION NUMBER: US/10/039,785  
 ; PRIOR FILING DATE: 2002-05-07  
 ; PRIOR APPLICATION NUMBER: 60/369,860  
 ; PRIOR FILING DATE: 2002-04-05  
 ; PRIOR APPLICATION NUMBER: 60/341,237  
 ; PRIOR FILING DATE: 2001-12-20  
 ; PRIOR APPLICATION NUMBER: 60/331,310  
 ; PRIOR FILING DATE: 2001-11-14  
 ; PRIOR APPLICATION NUMBER: 60/331,044  
 ; PRIOR FILING DATE: 2001-11-07  
 ; PRIOR APPLICATION NUMBER: 60/327,364  
 ; PRIOR FILING DATE: 2001-10-09  
 ; PRIOR APPLICATION NUMBER: 60/323,807  
 ; PRIOR FILING DATE: 2001-09-21  
 ; PRIOR APPLICATION NUMBER: 60/309,176  
 ; PRIOR FILING DATE: 2001-08-02  
 ; PRIOR APPLICATION NUMBER: 60/294,981  
 ; PRIOR FILING DATE: 2001-06-04  
 ; PRIOR APPLICATION NUMBER: 60/293,473  
 ; NUMBER OF SEQ ID NOS: 66  
 ; SOFTWARE: PatentIn Ver. 2.1  
 ; SEQ ID NO 48

TYPE: PRT  
 ORGANISM: Artificial sequence  
 FEATURE:  
 OTHER INFORMATION: T1014F11 scFv  
 US-10-039-785-48  
 Query Match 33.4%; Score 179.5; DB 4; Length 245;  
 Best Local Similarity 45.9%; Pred. No. 3.2e-12;  
 Matches 39; Conservative 13; Mismatches 26; Indels 7; Gaps 3;  
 QY 8 PGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAK 67  
 DB 148 PQQSVTISCTGTSDDVGXYK-VSWYQQHPGKAPKLMY----EVSMPSGVDPDRFSGSK 202  
 QY 68 DEAHNACVLITISVPQEDDADYCVS 92  
 DB 203 --SGNTASLTVSGIQAEDADYCA 225

RESULT 12  
 US-10-039-785-49  
 ; Sequence 49, Application US/10039785  
 ; Patent No. 6538938  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Salcedo et al.  
 ; TITLE OF INVENTION: Antibodies that Immunospecifically Bind to TRAIL  
 ; FILE REFERENCE: PF550  
 ; CURRENT APPLICATION NUMBER: US/10/039,785  
 ; PRIOR FILING DATE: 2002-05-07  
 ; PRIOR APPLICATION NUMBER: 60/369,860  
 ; PRIOR FILING DATE: 2002-04-05  
 ; PRIOR APPLICATION NUMBER: 60/341,237  
 ; PRIOR FILING DATE: 2001-12-20  
 ; PRIOR APPLICATION NUMBER: 60/331,310  
 ; PRIOR FILING DATE: 2001-11-14  
 ; PRIOR APPLICATION NUMBER: 60/331,044  
 ; PRIOR FILING DATE: 2001-11-07  
 ; PRIOR APPLICATION NUMBER: 60/327,364  
 ; PRIOR FILING DATE: 2001-10-09  
 ; PRIOR APPLICATION NUMBER: 60/323,807  
 ; PRIOR FILING DATE: 2001-09-21  
 ; PRIOR APPLICATION NUMBER: 60/309,176  
 ; PRIOR FILING DATE: 2001-08-02  
 ; PRIOR APPLICATION NUMBER: 60/294,981  
 ; PRIOR FILING DATE: 2001-06-04  
 ; PRIOR APPLICATION NUMBER: 60/293,473  
 ; NUMBER OF SEQ ID NOS: 66  
 ; SOFTWARE: PatentIn Ver. 2.1  
 ; SEQ ID NO 49

TYPE: PRT  
 ORGANISM: Artificial sequence  
 FEATURE:  
 OTHER INFORMATION: T1014G04 scFv  
 US-10-039-785-49  
 Query Match 33.2%; Score 178.5; DB 4; Length 245;  
 Best Local Similarity 44.7%; Pred. No. 4.1e-12;  
 Matches 38; Conservative 15; Mismatches 25; Indels 7; Gaps 3;  
 QY 8 PGQVAQLSCTLSPOHVTIRDYGVSWYQQRAGSAPRYLLYRSEEDHHRPADIPDRFSAK 67  
 DB 148 PQQSVTISCTGTSDDVGSYEY-VSWYQQHPGKAPRLMI----SEVNKRPSGVNRFSGSK 202  
 QY 68 DEAHNACVLITISVPQEDDADYCVS 92  
 DB 203 --SGNTASLTVSGIQAEDADYCVS 225

RESULT 13  
 US-09-240-274-68  
 ; Sequence 68, Application US/09240274  
 ; Patent No. 6255455  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Siegel, Donald L.  
 ; TITLE OF INVENTION: RH(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL  
 ; FILE REFERENCE: 09596-42U2  
 ; CURRENT APPLICATION NUMBER: US/09/240,274  
 ; PRIOR FILING DATE: 1999-01-29  
 ; PRIOR APPLICATION NUMBER: 60/081,380  
 ; PRIOR FILING DATE: 1998-04-10  
 ; PRIOR APPLICATION NUMBER: 60/028,550  
 ; PRIOR FILING DATE: 1996-10-11  
 ; NUMBER OF SEQ ID NOS: 224  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO 68  
 ; LENGTH: 108





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